
On the Alignment Between Supervised and Self-Supervised Contrastive Learning

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Abstract

Self-supervised contrastive learning (CL) has achieved remarkable empirical success, often producing representations that rival supervised pre-training on downstream tasks. Recent theory explains this by showing that the CL loss closely approximates a supervised surrogate, Negatives-Only Supervised Contrastive Learning (NSCL) loss, as the number of classes grows. Yet this loss-level similarity leaves an open question: *Do CL and NSCL also remain aligned at the representation level throughout training, not just in their objectives?*

We address this by analyzing the representation alignment of CL and NSCL models trained under shared randomness (same initialization, batches, and augmentations). First, we show that their induced representations remain similar: specifically, we prove that the similarity matrices of CL and NSCL stay close under realistic conditions. Our bounds provide high-probability guarantees on alignment metrics such as centered kernel alignment (CKA) and representational similarity analysis (RSA), and they clarify how alignment improves with more classes, higher temperatures, and its dependence on batch size. In contrast, we demonstrate that parameter-space coupling is inherently unstable: divergence between CL and NSCL weights can grow exponentially with training time.

Finally, we validate these predictions empirically, showing that CL–NSCL alignment strengthens with scale and temperature, and that NSCL tracks CL more closely than other supervised objectives. This positions NSCL as a principled bridge between self-supervised and supervised learning. Our code and project page are available at [code, project page].

1 Introduction

Self-supervised learning (SSL) has become the dominant approach for extracting transferable representations from large-scale unlabeled data. By leveraging training signals derived directly from the data, SSL methods avoid costly annotation while producing features that generalize across modalities, from vision (Chen et al., 2020; He et al., 2020; Zbontar et al., 2021; He et al., 2022; Oquab et al., 2024) to language (Gao et al., 2021; Reimers & Gurevych, 2019), speech (Schneider et al., 2019; Baevski et al., 2020; Hsu et al., 2021; Baevski et al., 2022), and vision–language (Radford et al., 2021; Jia et al., 2021; Zhai et al., 2023; Tschanne et al., 2025). Among SSL approaches, *contrastive learning* (CL) has been particularly successful: methods such as SimCLR (Chen et al., 2020), MoCo (He et al., 2020; Chen et al., 2021b), and CPC (van den Oord et al., 2019) train encoders by pulling together augmented views of the same input while pushing apart other samples. This simple principle has yielded state-of-the-art performance, often rivaling or surpassing supervised pre-training.

Despite this empirical success, a central puzzle remains: why does CL recover features so well aligned with semantic class boundaries? CL models often support nearly supervised-level downstream performance (Amir et al., 2022; Ben-Shaul et al., 2023; Weng et al., 2025), suggesting that supervision is somehow implicit in the objective. Recent theoretical progress sheds light on this: Luthra et al. (2025) showed that the CL objective closely approximates a supervised variant, *Negatives-Only Supervised Contrastive Learning (NSCL)*, where same-class samples are excluded from the denominator. Their analysis established that the CL–NSCL *losses* converge as the number of classes grows, and further characterized the geometry of NSCL minimizers and their linear probe performance. These results indicate that CL implicitly carries a supervised signal at the *loss level*.

Yet this view leaves a crucial question unresolved:

Do contrastive and supervised contrastive models remain aligned throughout training, not just at the level of their objectives?

Loss-level similarity does not guarantee that optimization paths coincide. In principle, differences in curvature, gradient noise, or learning rate schedules could amplify small loss discrepancies, causing stochastic gradient descent (SGD) trajectories to diverge. Thus, it remains unclear whether CL merely converges to a solution *similar* to NSCL, or whether their parameter and representations remain coupled across training.

Our contributions. We study the alignment between CL and NSCL under shared randomness (same initialization, mini-batches, and augmentations):

- **From drift to metrics.** The similarity control yields explicit, high-probability *lower bounds* on linear CKA and RSA at every epoch, showing that CL and NSCL representations remain non-trivially aligned and that the certified alignment tightens as C and τ grow, and it may increase with B depending on learning-rate scaling (Cors. 1–2). For completeness, we also bound parameter drift under β -smoothness (Thm. 2), which can grow exponentially even when representations remain aligned.
- **Empirical validation.** We validate our theory with experiments on CIFAR-10/100, Tiny-ImageNet, mini-ImageNet, and ImageNet-1K. We find that (i) CL–NSCL alignment strengthens with more classes and higher temperatures as well as correlates with the bound’s dependence on the batch size; and (ii) NSCL aligns with CL more strongly than other supervised learning methods (such as cross-entropy minimization and supervised contrastive learning (SCL) (Khosla et al., 2020)).

2 Related Work

A large body of work has sought to explain the success of contrastive learning (CL) from different perspectives. Early accounts linked CL to mutual information maximization between views of the same input (Bachman et al., 2019), though subsequent analyses showed that enforcing mutual information constraints too strongly can degrade downstream performance (McAllester & Stratos, 2020; Tschannen et al., 2020). A different line of work formalizes CL in terms of *alignment* and *uniformity* properties of the representation space (Wang & Isola, 2020; Wang & Liu, 2021; Chen et al., 2021a), capturing how positives concentrate while negatives spread across the sphere. These geometric criteria, while intuitive, do not fully explain how samples from different semantic classes are organized under CL training.

To address this, several papers have studied the ability of CL to recover latent clusters and semantic structures (Arora et al., 2019; Tosh et al., 2021; Zimmermann et al., 2021; Ash et al., 2022; Nozawa & Sato, 2021; HaoChen et al., 2021, 2022; Shen et al., 2022; Wang et al., 2022; Awasthi et al., 2022; Bao et al., 2022). Most of these results rely on restrictive assumptions, such as conditional independence of augmentations given cluster identity (Arora et al., 2019; Tosh et al., 2021; Saunshi et al., 2022; Awasthi et al., 2022). To weaken such assumptions, HaoChen & Ma (2023) proposed analyzing spectral contrastive objectives that encourage cluster preservation without requiring augmentation connectivity, while Parulekar et al. (2023) showed that InfoNCE itself learns cluster-preserving embeddings when the hypothesis class is capacity-limited.

Another perspective comes from linking CL to supervised learning. For instance, Balestrieri & LeCun (2024) showed that in linear models, self-supervised objectives such as VicReg coincide with supervised quadratic losses. Building on this supervised view, Luthra et al. (2025) established an explicit coupling between the InfoNCE contrastive loss and a supervised variant that removes positives from the denominator. In contrast to prior results, these bounds are label-agnostic, architecture-independent, and hold uniformly throughout optimization.

Beyond clustering and supervision, other theoretical studies have examined different aspects of CL: feature learning dynamics in linear and shallow nonlinear networks (Tian, 2022; Ji et al., 2023; Wen & Li, 2021; Tian, 2023), the role and optimality of augmentations (Tian et al., 2020; Feigin et al., 2025), the projection head (Gupta et al., 2022; Gui et al., 2023; Xue et al., 2024; Ouyang et al., 2025), sample complexity (Alon et al., 2024), and strategies to reduce batch-size requirements (Yuan et al., 2022). Finally, several works explore connections between contrastive and non-contrastive SSL paradigms (Wei et al., 2021; Balestrieri & LeCun, 2022; Lee et al., 2021; Garrido et al., 2023; Shwartz-Ziv et al., 2023).

3 Problem Setup

We work with a class-balanced dataset $S = \{(x_i, y_i)\}_{i=1}^N \subset \mathcal{X} \times [C]$, where $[C] = \{1, \dots, C\}$ and each class c contributes n examples ($N = Cn$). An encoder $f_w : \mathcal{X} \rightarrow \mathbb{R}^d$ with parameters $w \in \mathbb{R}^p$ maps inputs to embeddings. Similarity is measured by a bounded function $\text{sim} : \mathbb{R}^d \times \mathbb{R}^d \rightarrow [-1, 1]$; throughout our experiments we use cosine similarity on ℓ_2 -normalized embeddings, $\text{sim}(u, v) = \langle u, v \rangle / (\|u\| \|v\|)$, which satisfies this bound.

Data augmentations are modeled by a Markov kernel $\alpha(\cdot | x)$ on \mathcal{X} : given x , we draw an independent view $x' \sim \alpha(x)$. Unless stated otherwise, augmentation draws are independent across samples, across repeated views of the same sample, and across training steps. We write $x'_i \sim \alpha(x_i)$ for a single view and $(x_i^{(1)}, x_i^{(2)}) \stackrel{\text{i.i.d.}}{\sim} \alpha(x_i)$ for two views of the same input.

Fix a batch size $B \in \mathbb{N}$. A batch is a multiset $\mathcal{B} = \{(x_{j_t}, x'_{j_t}, y_{j_t})\}_{t=1}^B$ sampled with replacement from S , together with independent augmentations $x'_{j_t} \sim \alpha(x_{j_t})$. For any anchor triplet $(x_i, x'_i, y_i) \in \mathcal{B}$, define the *per-anchor CL loss* and the *CL batch loss* as:

$$\begin{aligned} \ell_{i, \mathcal{B}}(w) &:= -\log \left(\frac{\exp(\text{sim}(f_w(x_i), f_w(x'_i)) / \tau)}{\sum_{(x_j, x'_j, y_j) \in \mathcal{B}} [\mathbf{1}\{j \neq i\} \exp(\text{sim}(f_w(x_i), f_w(x_{j_t})) / \tau) + \exp(\text{sim}(f_w(x_i), f_w(x'_{j_t})) / \tau)]} \right), \\ \bar{\ell}_{\mathcal{B}}^{\text{CL}}(w) &:= \frac{1}{B} \sum_{(x_i, x'_i, y_i) \in \mathcal{B}} \ell_{i, \mathcal{B}}(w). \end{aligned}$$

For the same realized batch \mathcal{B} , define the per-anchor *negative subset* $\mathcal{B}_i^- := \{(x_{j_t}, x'_{j_t}, y_{j_t}) \in \mathcal{B} : y_{j_t} \neq y_i\}$. The *NSCL per-anchor* and *batch* losses are

$$\begin{aligned} \ell_{i, \mathcal{B}_i^-}(w) &:= -\log \left(\frac{\exp(\text{sim}(f_w(x_i), f_w(x'_i)) / \tau)}{\sum_{(x_j, x'_j, y_j) \in \mathcal{B}_i^-} [\exp(\text{sim}(f_w(x_i), f_w(x_j)) / \tau) + \exp(\text{sim}(f_w(x_i), f_w(x'_j)) / \tau)]} \right), \\ \bar{\ell}_{\mathcal{B}}^{\text{NSCL}}(w) &:= \frac{1}{B} \sum_{(x_i, x'_i, y_i) \in \mathcal{B}} \ell_{i, \mathcal{B}_i^-}(w). \end{aligned}$$

Prior work (Luthra et al., 2025) shows that the CL–NSCL loss gap is uniformly $\mathcal{O}(1/C)$, but what we ultimately care about is whether the *embeddings* align. To quantify representation similarity we use linear Centered Kernel Alignment (CKA) (Kornblith et al., 2019) and Representation Similarity Analysis (RSA) (Kriegeskorte et al., 2008) defined on cosine-similarity matrices: for N common inputs with embeddings $Z = \{z_i\}_{i=1}^N$ and $Z' = \{z'_i\}_{i=1}^N$, let $\Sigma(Z)_{ij} = \cos(z_i, z_j)$ and $H = I - \frac{1}{N}\mathbf{1}\mathbf{1}^\top$; linear CKA is

$$\text{CKA}(Z, Z') = \frac{\langle H\Sigma(Z)H, H\Sigma(Z')H \rangle_F}{\|H\Sigma(Z)H\|_F \|H\Sigma(Z')H\|_F},$$

and RSA is the *Pearson* correlation between the (upper-triangular) off-diagonal entries of the dissimilarity matrices $\text{RDM}(Z) = \mathbf{1}\mathbf{1}^\top - \Sigma(Z)$ and $\text{RDM}(Z') = \mathbf{1}\mathbf{1}^\top - \Sigma(Z')$:

$$\text{RSA}(Z, Z') = \text{Corr}(\text{vec}_\Delta(\text{RDM}(Z)), \text{vec}_\Delta(\text{RDM}(Z'))),$$

where vec_Δ stacks the upper-triangular entries ($i < j$) column-wise.

This raises the following question: *Beyond a small objective gap, does training CL and NSCL actually lead to similar representations (e.g., high CKA/RSA)?*

In the spirit of Thm. 1 of Luthra et al. (2025), we prove that when two runs use shared randomness (same initialization, mini-batches, and augmentations), the per-step gradient mismatch is uniformly bounded (Lem. 7). Similarly, we show that the CL and NSCL similarity matrices remain close throughout training (Thm. 1), which yields explicit CKA/RSA lower bounds (Cors. 1-2).

Additional notation for high-probability factors. Fix a training horizon $T \in \mathbb{N}$, a confidence level $\delta \in (0, 1)$, and a temperature $\tau > 0$. For later use, define $\epsilon_{B,\delta} := \sqrt{\frac{1}{2B} \log\left(\frac{TB}{\delta}\right)}$ and $\Delta_{C,\delta}(B; \tau) := \frac{2e^{2/\tau} \left(\frac{1}{C} + \epsilon_{B,\delta}\right)}{1 - \frac{1}{C} - \epsilon_{B,\delta}}$, and assume $\epsilon_{B,\delta} < 1 - \frac{1}{C}$ so the denominator is positive.

4 Theory

We examine how contrastive learning (CL) and negatives-only supervised contrastive learning (NSCL) co-evolve when initialized identically and trained with the same mini-batches and augmentations. While one might first attempt to study their trajectories in parameter space, such an approach quickly breaks down: without strong assumptions on the loss landscape (e.g., convexity or strong convexity), small reparameterizations can distort distances, and nonconvex dynamics cause parameter drift to grow uncontrollably over time (see App. B). For this reason, we set weight-space coupling aside and turn instead to the aspect that directly shapes downstream behavior—the *representations*—analyzing their alignment in similarity space.

4.1 Coupling in Representation (Similarity) Space

Let $\Sigma_t \in [-1, 1]^{N \times N}$ denote the pairwise similarity matrix of a fixed reference set at step t (cosine similarity of normalized embeddings; diagonals are 1). We analyze the coupled evolution of the CL and NSCL similarities,

$$\Sigma_t^{\text{CL}}, \Sigma_t^{\text{NSCL}} \in [-1, 1]^{N \times N},$$

under identical mini-batches and augmentations. This representation-space view is invariant to reparameterization and directly tracks representational geometry.

Surrogate similarity dynamics. To make the analysis explicit, we work with a “similarity-descent” surrogate that updates only those entries touched by the current batch. For a realized mini-batch $\mathcal{B}_t = \{(x_j, x'_j, y_j)\}_{j=1}^B$ (with $x'_j \sim \alpha(x_j)$), let $\bar{\ell}_{\mathcal{B}_t}^{\text{CL}}(\Sigma)$ and $\bar{\ell}_{\mathcal{B}_t}^{\text{NSCL}}(\Sigma)$ be the usual InfoNCE-type losses written as functions of the relevant similarity entries (with temperature $\tau > 0$). Define the batch-gradient maps

$$G_t^{\text{CL}} := \nabla_\Sigma \bar{\ell}_{\mathcal{B}_t}^{\text{CL}}(\Sigma_t^{\text{CL}}), \quad G_t^{\text{NSCL}} := \nabla_\Sigma \bar{\ell}_{\mathcal{B}_t}^{\text{NSCL}}(\Sigma_t^{\text{NSCL}}),$$

setting all untouched entries to zero. The surrogate updates are

$$\Sigma_{t+1}^{\text{CL}} = \Sigma_t^{\text{CL}} - \eta_t G_t^{\text{CL}}, \quad \Sigma_{t+1}^{\text{NSCL}} = \Sigma_t^{\text{NSCL}} - \eta_t G_t^{\text{NSCL}}, \quad (1)$$

with shared initialization and shared randomness (same \mathcal{B}_t and augmentations). In App. C we show that these surrogate dynamics faithfully track the similarity evolution induced by parameter-space SGD. We now formalize the coupling bound.

Theorem 1 (Similarity-space coupling). *Fix $B, T \in \mathbb{N}$, $\delta \in (0, 1)$, and temperature $\tau > 0$. Consider the coupled similarity-descent recursions equation 1 for CL and NSCL with shared initialization and shared mini-batches/augmentations. Then, with probability at least $1 - \delta$ over the draws of the mini-batches and augmentations, for any stepsizes $(\eta_t)_{t=0}^{T-1}$,*

$$\|\Sigma_T^{\text{CL}} - \Sigma_T^{\text{NSCL}}\|_F \leq \exp\left(\frac{1}{2\tau^2 B} \sum_{t=0}^{T-1} \eta_t\right) \frac{1}{\tau\sqrt{B}} \left(\sum_{t=0}^{T-1} \eta_t\right) \Delta_{C,\delta}(B; \tau). \quad (2)$$

From similarity drift to CKA/RSA guarantees. We translate the high-probability control on the similarity drift from Thm. 1, into bounds on two standard representational metrics.

CKA. Recall from Sec. 3 that linear CKA (Kornblith et al., 2019) is the normalized Frobenius inner product between centered similarity matrices. $H := I - \frac{1}{N}\mathbf{1}\mathbf{1}^\top$ be the centering projector and define centered Gram matrices $K_T^{\text{CL}} := H\Sigma_T^{\text{CL}}H$ and $K_T^{\text{NSCL}} := H\Sigma_T^{\text{NSCL}}H$. The (linear) CKA at step T is $\text{CKA}_T = \frac{\langle K_T^{\text{CL}}, K_T^{\text{NSCL}} \rangle_F}{\|K_T^{\text{CL}}\|_F \|K_T^{\text{NSCL}}\|_F} \in [0, 1]$. Because $\|H X H\|_F \leq \|X\|_F$, any bound on $\|\Sigma_T^{\text{CL}} - \Sigma_T^{\text{NSCL}}\|_F$ controls $\|K_T^{\text{CL}} - K_T^{\text{NSCL}}\|_F$. For convenience, introduce the relative deviation $\rho_T := \frac{\|K_T^{\text{CL}} - K_T^{\text{NSCL}}\|_F}{\|K_T^{\text{CL}}\|_F}$.

Corollary 1 (CKA lower bound). *In the setting of Thm. 1. Assume $\|K_T^{\text{CL}}\|_F > 0$. With probability at least $1 - \delta$,*

$$\text{CKA}_T \geq \frac{1 - \rho_T}{1 + \rho_T}, \quad \rho_T \leq \frac{\exp\left(\frac{1}{2\tau^2 B} \sum_{t=0}^{T-1} \eta_t\right) \frac{1}{\tau\sqrt{B}} \left(\sum_{t=0}^{T-1} \eta_t\right) \Delta_{C,\delta}(B; \tau)}{\|K_T^{\text{CL}}\|_F}.$$

RSA. Recall from Sec. 3 that RSA (Kriegeskorte et al., 2008) is the Pearson correlation between the off-diagonal entries of representational dissimilarity matrices (RDMs). Let $M = \binom{N}{2}$ and define off-diagonal RDM vectors $a_T, b_T \in \mathbb{R}^M$ by $a_T(u, v) = 1 - \Sigma_T^{\text{CL}}(u, v)$ and $b_T(u, v) = 1 - \Sigma_T^{\text{NSCL}}(u, v)$ for $u < v$. Write $\sigma_{D,T} > 0$ for the empirical standard deviation of the entries of a_T . The RSA score is the Pearson correlation $\text{RSA}_T = \text{Corr}(a_T, b_T)$. Zeroing the diagonal does not increase Frobenius norms, so $\|b_T - a_T\|_2 \leq \|\Sigma_T^{\text{NSCL}} - \Sigma_T^{\text{CL}}\|_F$. It will be useful to measure the relative discrepancy $r_T := \frac{\|b_T - a_T\|_2}{\sqrt{M} \sigma_{D,T}}$.

Corollary 2 (RSA lower bound). *In the setting of Thm. 1. Assume $\sigma_{D,T} > 0$. With probability at least $1 - \delta$,*

$$\text{RSA}_T \geq \frac{1 - r_T}{1 + r_T}, \quad r_T \leq \frac{\exp\left(\frac{1}{2\tau^2 B} \sum_{t=0}^{T-1} \eta_t\right) \frac{1}{\tau\sqrt{B}} \left(\sum_{t=0}^{T-1} \eta_t\right) \Delta_{C,\delta}(B; \tau)}{\sqrt{M} \sigma_{D,T}}.$$

These results complement the parameter-space analysis. While parameter trajectories may diverge exponentially (in the non-convex setting; see App. D.3), the induced similarities—and hence representational metrics such as CKA and RSA—remain tightly controlled by class count, batch size, learning rate, and temperature τ . The key quantity is the similarity-matrix drift $\|\Sigma_T^{\text{CL}} - \Sigma_T^{\text{NSCL}}\|_F$, which Thm. 1 bounds in two stabilizing ways.

First, the exponential factor is moderated by the $\frac{1}{\tau^2 B}$ term in the exponent. Unlike parameter space, where the growth rate scales with β , the “instability rate” in similarity space is only $\frac{1}{2\tau^2 B}$ and is therefore negligible for typical batch sizes (e.g., $B \approx 10^2\text{--}10^3$).

Second, the prefactor $\frac{1}{\tau\sqrt{B}} \left(\sum_t \eta_t\right) \Delta_{C,\delta}(B; \tau)$ decreases rapidly with batch size and class count (note $\Delta_{C,\delta}(B; \tau)$ shrinks with larger C and grows with smaller τ through $e^{2/\tau}$). In practical regimes ($C \sim 10^3$, $B \sim 10^2\text{--}10^3$), this prefactor is small, making the total Frobenius gap negligible relative to the scale of the similarity matrices.

Together, these effects yield high-probability guarantees $\text{CKA}_T \geq (1 - \rho_T)/(1 + \rho_T)$ and $\text{RSA}_T \geq (1 - r_T)/(1 + r_T)$ with $\rho_T, r_T \ll 1$ in realistic conditions. Thus, even if parameters drift, the induced

representations evolve in a coupled and stable manner—consistent with empirical findings that CL and NSCL remain closely aligned in practice.

Proof idea. We begin with a high-probability batch-composition guarantee (Cor. 3): with probability at least $1 - \delta$, every anchor’s denominator contains the expected proportion of negatives up to an $\epsilon_{B,\delta}$ fluctuation. This rules out positive-heavy batches that would otherwise cause the NSCL renormalization to deviate substantially from CL. Conditioning on this event, the CL–NSCL batch-gradient gap decomposes into (i) a *reweighting error*, bounded in total variation by $\Delta_{C,\delta}(B; \tau)$ (Lem. 6), and (ii) a *stability term* from the dependence on the current similarities, controlled by the $\frac{1}{2\tau^2 B}$ -Lipschitzness of the batch-gradient map in Frobenius norm (Lem. 2 at temperature τ). Using block-orthogonality across anchors (Lem. 1), the reweighting contributions combine in quadrature, giving the per-step estimate (Lem. 8),

$$\|G_t^{\text{CL}}(\Sigma_t^{\text{CL}}) - G_t^{\text{NSCL}}(\Sigma_t^{\text{NSCL}})\|_F \leq \frac{1}{\tau} \cdot \frac{\Delta_{C,\delta}(B; \tau)}{\sqrt{B}} + \frac{1}{2\tau^2 B} \|\Sigma_t^{\text{CL}} - \Sigma_t^{\text{NSCL}}\|_F.$$

Consequently, the similarity drift satisfies the recurrence

$$\|\Sigma_{t+1}^{\text{CL}} - \Sigma_{t+1}^{\text{NSCL}}\|_F \leq \left(1 + \frac{\eta_t}{2\tau^2 B}\right) \|\Sigma_t^{\text{CL}} - \Sigma_t^{\text{NSCL}}\|_F + \eta_t \frac{1}{\tau} \cdot \frac{\Delta_{C,\delta}(B; \tau)}{\sqrt{B}},$$

where the first term propagates existing error and the second injects the new discrepancy introduced at step t . Unrolling this recurrence (discrete Grönwall) yields

$$\|\Sigma_T^{\text{CL}} - \Sigma_T^{\text{NSCL}}\|_F \leq \exp\left(\frac{1}{2\tau^2 B} \sum_{t=0}^{T-1} \eta_t\right) \frac{1}{\tau \sqrt{B}} \left(\sum_{t=0}^{T-1} \eta_t\right) \Delta_{C,\delta}(B; \tau).$$

Finally, centering contracts Frobenius norms, so this control transfers directly to the centered Gram matrices, and applying standard $(1 - \rho)/(1 + \rho)$ and $(1 - r)/(1 + r)$ comparisons yields the claimed CKA/RSA lower bounds.

5 Experiments

Datasets and augmentations. We experiment with the following standard vision classification datasets - CIFAR10 and CIFAR100 (Krizhevsky, 2009), Mini-ImageNet (Vinyals et al., 2016), Tiny-ImageNet (Han, 2020), and ImageNet-1K (Deng et al., 2009). (See App. A for details.)

Methods, architectures, and optimizers. For all our experiments, we have followed the SimCLR (Chen et al., 2020) algorithm. We use a ResNet-50 (He et al., 2016) encoder with a width-multiplier factor of 1. The projection head follows a standard two-layer MLP architecture composed of: Linear($2048 \rightarrow 2048$) \rightarrow ReLU \rightarrow Linear($2048 \rightarrow 128$). For cross-entropy training, we attach an additional classification head Linear($128 \rightarrow C$) where C is the number of classes.

For contrastive learning, we use the DCL loss that avoids positive-negative coupling during training (Yeh et al., 2022). For supervised learning, we use the following variants: Supervised Contrastive Loss (Khosla et al., 2020), Negatives-Only Supervised Contrastive Loss (Luthra et al., 2025), and Cross-Entropy Loss (Shannon, 1948). To minimize the loss, we adopt the LARS optimizer (You et al., 2017) which has been shown in (Chen et al., 2020) to be effective for training with large batch sizes. For LARS, we set the momentum to 0.9 and the weight decay to 10^{-6} . All experiments are carried out with a batch size of $B = 1024$. The base learning rate is scaled with batch size as $0.3 \cdot \lfloor B/256 \rfloor$, following standard practice (Chen et al., 2020). We employ a warm-up phase (Goyal et al., 2017) for the first 10 epochs, followed by a cosine learning rate schedule without restarts (Loshchilov & Hutter, 2016) for the remaining epochs. All models were trained on a single node with one 94 GB NVIDIA H100 GPUs.

Evaluation metrics. To quantitatively measure the alignment between the learned representation spaces of different models, we monitor linear CKA and RSA (check Sec. 3 for details) during training. Both CKA and RSA range from 0 to 1, where 1 indicates identical similarity structures. For completeness, along with CKA and RSA, we also report downstream performance via Nearest Class Center Classifier (Galanti et al., 2022) and Linear Probing (Köhn, 2015; Gupta et al., 2015; Belinkov, 2022) accuracies in Tab. 1.

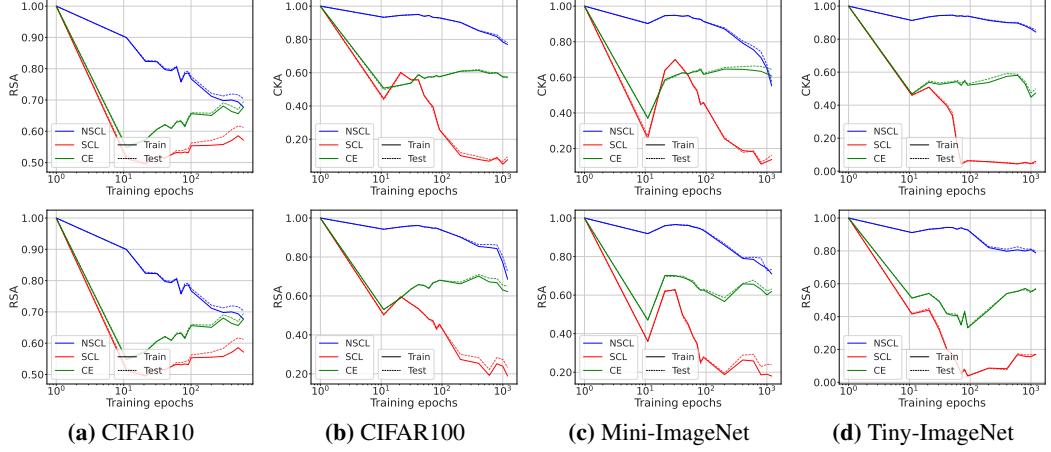


Figure 2: Alignment during training. We train ResNet-50 models with decoupled CL, SCL, NSCL, and CE. For the first 1,000 epochs, the CL-trained model is substantially more aligned with the NSCL-trained model than with the others. However, alignment declines when training continues much longer.

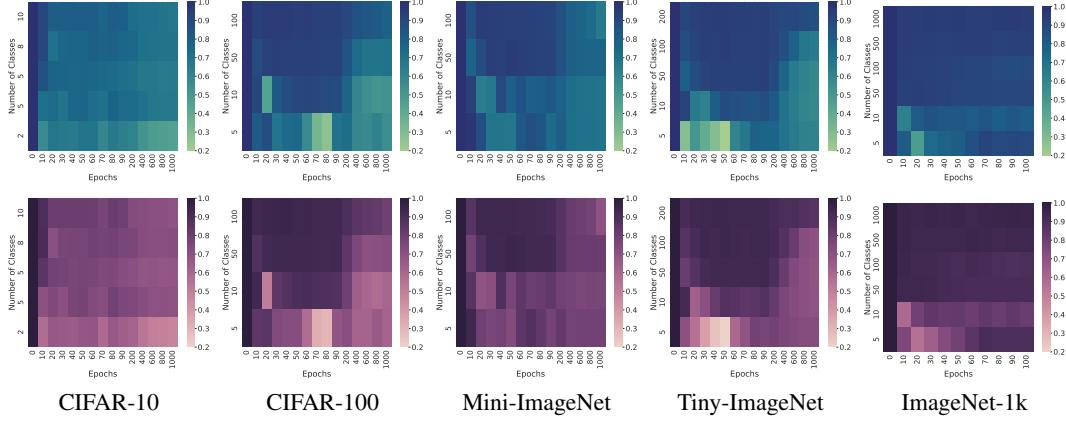


Figure 3: CL-NSCL alignment (linear CKA) increases with the number of training classes. The heatmaps show the linear CKA between CL and NSCL models. For each dataset, we visualize alignment on the training (top row, green) and test (bottom row, purple) sets. The y-axis indicates the number of classes (N) used for training, and the x-axis represents the training epoch. While alignment is consistently higher for larger N , it also tends to decrease as training progresses for any fixed N .

5.1 Experimental Results

Alignment analysis as a function of epochs. To understand how representational similarity evolves, we trained a model with a CL objective and monitored its alignment (via CKA/RSA) against supervised models trained with NSCL, CE, and SCL. We find that NSCL consistently achieves the highest alignment with CL throughout training across multiple datasets compared to CE and SCL (see Fig. 2). For example, after 1k epochs on Tiny-ImageNet, the CL-NSCL alignment reaches a CKA of 0.87, in contrast to just 0.043 for CL-SCL.

Intuitively, this behavior stems from the geometric constraints induced by the respective loss functions. NSCL’s objective structurally mimics CL’s. Both losses attract a single positive sample to an anchor for each instance, encouraging a similar geometric arrangement focused on instance-level discrimination. SCL enforces a much stronger, class-level constraint. By pulling all positive samples in a batch together, it aggressively reduces intra-class variance to form tight clusters. This induces a geometric structure different from the one learned by CL’s instance-based task (Luthra et al., 2025). Cross-Entropy (CE) represents an intermediate case. It focuses on class separability via a decision

	CIFAR-10		CIFAR-100		Mini-ImageNet		Tiny-ImageNet	
	NCCC	LP	NCCC	LP	NCCC	LP	NCCC	LP
CL	88.37	90.16	54.62	65.65	60.78	65.30	40.59	44.61
NSCL	94.47	94.09	60.14	68.38	63.92	72.60	40.76	45.79
SCL	94.93	94.67	64.06	69.52	74.78	76.00	48.63	48.73
CE	92.97	93.39	67.35	68.04	75.20	74.00	48.28	52.57

Table 1: Nearest Class-Center Classifier (NCCC) and Linear Probe (LP) test accuracies (%). We report the accuracies against the all-way classification task in each dataset. The models (also used in Fig. 2) were pre-trained on their respective datasets: CIFAR10, CIFAR100, Mini-ImageNet and Tiny-ImageNet.

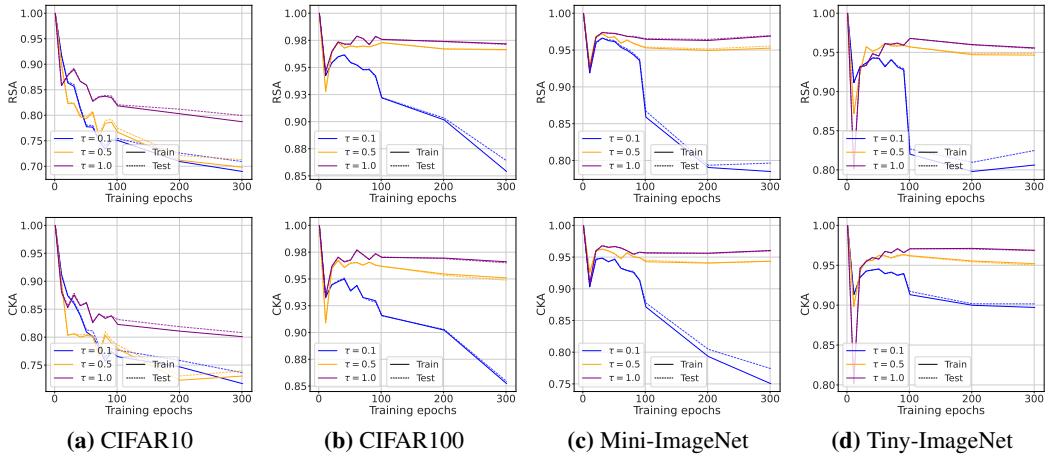


Figure 4: **Higher τ increases the CL-NSCL alignment.** The plots show RSA (top row) and CKA (bottom row) over 300 epochs. We trained CL and NSCL models with varying temperatures ($\tau \in \{0.1, 0.5, 1.0\}$) on four datasets. Across all datasets, a higher temperature $\tau = 1.0$ (shown in purple) evidently results in the highest alignment.

boundary rather than explicit feature clustering. This explains its modest alignment trajectory, which dips initially before rising as both CL and CE converge toward learning semantic features.

Validating Thm. 1 as a function of class count. Our theory (Thm. 1) predicts that using more classes yields stronger CL-NSCL alignment. We test this via C' -way training: for each $C' \in [2, C]$, we train CL and NSCL on random C' -class subsets for 1,000 epochs (except 100 epochs for IM-1K). As shown in Fig. 3, representation similarity (RSA/CKA) increases with C' across all datasets.

Effect of temperature on alignment. According to our theoretical analysis (Cors. 1–2), CL-NSCL alignment improves with higher values of temperature (τ). We empirically verify this claim by training CL and NSCL models for 300 epochs, over three different values of $\tau \in \{0.1, 0.5, 1.0\}$. Both models—CL and NSCL—are trained with same τ in each run. As shown in Fig. 4, models trained with $\tau = 1.0$ achieve higher alignment compared to models trained with lower temperatures.

Effect of batch size on alignment. Our framework also predicts that alignment varies with batch-size B depending on how the learning rate scales. To investigate this, we vary η with B across four cases: $\eta = \frac{0.3B}{256}$, $\eta = \frac{0.3\sqrt{B}}{256}$, $\eta = \frac{0.3\sqrt[4]{B}}{256}$, and $\eta = 0.3$. Under $\mathcal{O}(B)$ scaling, CL-NSCL alignment decreases as B grows, matching the theorem’s implication for that scaling; for the other three cases, alignment increases with B , again consistent with the bound under those dependencies (see Fig. 5).

Weight-space coupling. We next study whether the observed alignment between representations of contrastive and supervised models is also reflected directly in their parameters. For this, we measure the average weight difference between a contrastive model and two supervised counterparts as follows: $\sum_l \frac{\|w_{\text{CL}}^l - w_{\text{sup}}^l\|_F}{0.5 (\|w_{\text{CL}}^l\|_F + \|w_{\text{sup}}^l\|_F)}$ where w_{CL}^l and w_{sup}^l are weights corresponding to l^{th} layer of

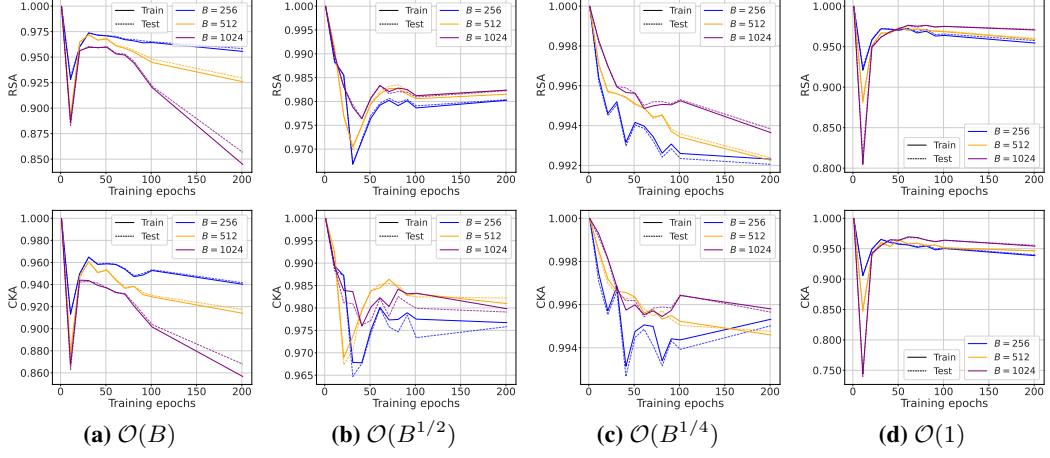


Figure 5: Effect of batch size with scaled learning rates. We trained CL, and NSCL models for 300 epochs on Mini-ImageNet, with varying batch-sizes ($B \in \{256, 512, 1024\}$). For each experiment, the learning rate η is scaled as a function of batch-size, as mentioned under each panel. For instance, the results shown in panel (b) use a learning rate of $\eta = \frac{0.3\sqrt{B}}{256}$.

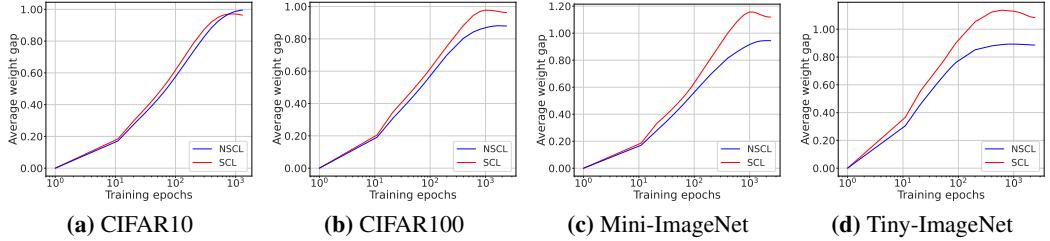


Figure 6: Weight-space alignment quickly deteriorates. Using the same ResNet-50 instances as in Fig. 2, we plot the average weight gap between CL and the supervised models (NSCL and SCL) across training epochs. Both supervised variants diverge from the CL model, with SCL showing a wider separation.

self-supervised and supervised models respectively, and $\|\cdot\|_F$ denotes Frobenius norm. As we show in Fig. 6, for each dataset, we observe a significant divergence in weight space: both supervised models (NSCL and SCL) increasingly separate from the contrastive model as training progresses.

6 Conclusions, Limitations and Future Work

Conclusions. We studied the dynamic alignment between contrastive learning (CL) and its supervised counterpart (NSCL). By analyzing coupled SGD under shared randomness, we showed that while parameter-space trajectories may diverge exponentially, representation-space dynamics are far more stable: the similarity matrices induced by CL and NSCL remain close throughout training. This yields high-probability lower bounds on alignment metrics such as CKA and RSA, directly certifying representational coupling. Empirically, our experiments confirmed these trends across datasets and architectures. Together, our results highlight that the implicit supervised signal in CL is not confined to its loss function but extends throughout the entire optimization trajectory.

Limitations. Our theoretical bounds, while structurally informative, are quantitatively loose in large-scale, long-horizon regimes. The exponential factor can quickly make the bounds vacuous beyond the first few epochs of large-scale training. This limitation stems from worst-case arguments and uniform high-probability concentration bounds that favor generality over sharpness. Nonetheless, the analysis offers qualitative insights into how the temperature parameter, the number of hidden classes, the batch size and training duration affect the alignment between CL and NSCL models.

Future directions. We view our results as a first step toward a more refined theory of self-supervised representation alignment. Future work could (i) derive tighter constants by exploiting data-dependent structure rather than worst-case bounds, and (ii) extend the framework to other SSL paradigms (e.g., non-contrastive methods). Improving these guarantees while retaining their stability properties would provide an even stronger theoretical bridge between supervised and self-supervised learning.

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A Additional Experiments

Datasets and augmentations. CIFAR10 and CIFAR100 both consist of 50000 training images and 10000 validation images with 10 classes and 100 classes, respectively, uniformly distributed across the dataset, i.e., CIFAR10 has 5000 samples per class and CIFAR100 has 500 samples per class. Mini-ImageNet also has 5000 test images on top of 50000 train and 10000 validation images, with 100 of 1000 classes from ImageNet-1K (Deng et al., 2009) (at the original resolution). Tiny-ImageNet contains 100000 images downsampled to 64×64 , with total 200 classes from IM-1K. Each class has 500 training, 50 validation, and 50 test images.

We use standard augmentations as proposed in SimCLR (Chen et al., 2020). For experiments on Mini-ImageNet, we use the following pipeline: random resized cropping to 224×224 , random horizontal flipping, color jittering (brightness, contrast, saturation: 0.8; hue: 0.2), random grayscale conversion ($p = 0.2$), and Gaussian blur (applied with probability 0.1 using a 3×3 kernel and $\sigma = 1.5$). For Tiny-ImageNet, we drop saturation to 0.4 and hue to 0.1 due to low resolution images. For CIFAR datasets, we adopt a similar pipeline with appropriately scaled parameters. The crop size is adjusted to 32×32 , and the color jitter parameters are scaled to saturation 0.4, and hue 0.1.

A.1 Effect of number of classes on alignment

In addition to the linear CKA results reported in the main text (Fig. 3), we also evaluate representational similarity using RSA. The corresponding RSA values are presented in Fig. 7, providing a complementary perspective on alignment across varying numbers of classes.

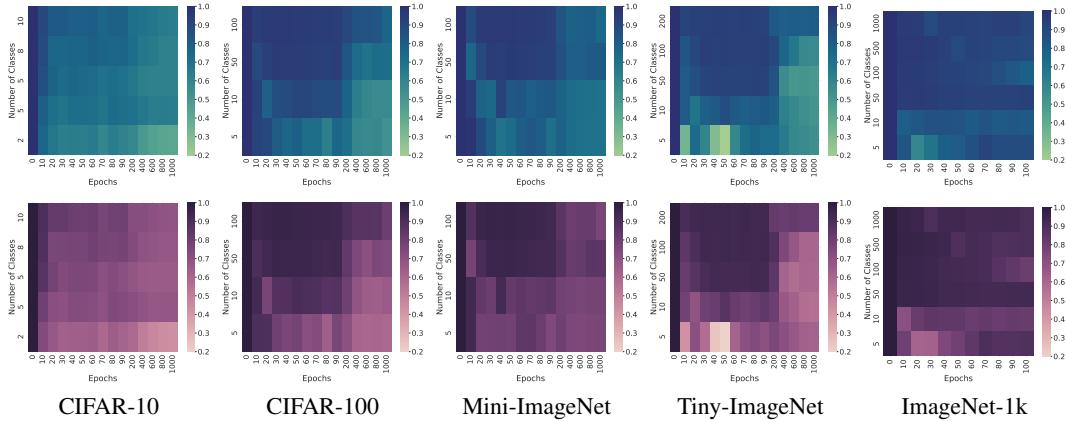


Figure 7: **CL-NSCL alignment (RSA) increases with the number of training classes.** See Sec. 5.1 and Fig. 3 for experimental details.

A.2 Fig. 1 Methodology

We explain how to generate the plots comparing alignment in weight-space and representation-space. The two plots on the left visualize the direction of learning for each model. Each vector represents the change in model’s state from initialization (epoch 0) to epoch 1000.

Model states. We consider CL and NSCL models trained on CIFAR100, corresponding to epoch 0 and epoch 1000-a total of four models.

Weight space. This plot shows how the raw parameters evolve during training. For all four models, we first flatten all the weights into a massive vector which gives us four points in a very high dimensional space (order of 10^7). To visualize these points, we perform Principal Component Analysis (PCA) on all four vectors combined and fit them to a 3D space. This creates a shared 3D coordinate system. We transform all four points into this space and we get $p_{\text{CL}}^0, p_{\text{CL}}^{1000}, p_{\text{NSCL}}^0, p_{\text{NSCL}}^{1000}$. Using these points, we create two vectors: $(v_{\text{CL}}, v_{\text{NSCL}})$, and create polar plot using the final vectors and the calculated angle between them (85.7°).

Representation space. This plot shows how model’s alignment for a specific class evolved. We pick one class from our dataset (CIFAR100) and randomly sample 100 images. We use the same

samples for all four models to extract their corresponding features, say $Z \in \mathbb{R}^{100 \times d}$, where d is the projection dimension. We concatenate total 400 representations (100 from each model) and perform PCA to learn a shared 3D coordinate system. The representations are transformed to this shared space ($\mathbb{R}^{100 \times d} \rightarrow \mathbb{R}^{100 \times 3}$) and averaged to a single 3D point for each model. Just like before, a polar plot is created using the vectors and angle between them (27.8°).

Similarity metrics. We report RSA and CKA values computed between DCL and NSCL models trained on CIFAR100. Additionally, we show their average weight gap as detailed in Sec. 5.1. It is evident that models stay aligned in representation space but diverge in weight space.

B Parameter-Space Coupling

To complement the analysis in Sec. 4, we compare the two trajectories in parameter space. Let $e_t = \|w_t^{\text{CL}} - w_t^{\text{NSCL}}\|$ denote the parameter drift at step t . We would like to bound it as a function of the number of training iterations, batch size, and learning rate scheduling. We use classic techniques that can be found at (Bousquet & Elisseeff, 2002; Hardt et al., 2016; Mou et al., 2018; Kuzborskij & Lampert, 2017).

Optimization. In order to isolate the effect of the loss, we optimize both objectives (CL and NSCL) with standard mini-batch SGD under a single coupled protocol: at step t we draw a batch $\mathcal{B}_t = \{(x_j, x'_j, y_j)\}_{j=1}^B$ with replacement, where each $x'_j \sim \alpha(x_j)$ (e.g., random crop/resize, horizontal flip, color jitter, Gaussian blur); we average per-anchor terms to form either $\bar{\ell}_{\mathcal{B}_t}^{\text{CL}}(w)$ or $\bar{\ell}_{\mathcal{B}_t}^{\text{NS}}(w)$ using cosine similarity (optionally temperature-scaled), hence bounded in $[-1, 1]$; and we update $w_{t+1} = w_t - \eta_t \nabla \bar{\ell}_{\mathcal{B}_t}(w_t)$ with prescribed $\eta_t > 0$. We then run two coupled SGD trajectories from the same initialization $w_0^{\text{CL}} = w_0^{\text{NSCL}}$ that share the *same* batches and augmentations $(\mathcal{B}_t)_{t=0}^{T-1}$ and differ only by NSCL’s exclusion of same-class negatives:

$$w_{t+1}^{\text{CL}} = w_t^{\text{CL}} - \eta_t \nabla \bar{\ell}_{\mathcal{B}_t}^{\text{CL}}(w_t^{\text{CL}}), \quad w_{t+1}^{\text{NSCL}} = w_t^{\text{NSCL}} - \eta_t \nabla \bar{\ell}_{\mathcal{B}_t}^{\text{NSCL}}(w_t^{\text{NSCL}}), \quad t = 0, \dots, T-1.$$

Throughout the analysis, we make standard assumptions on the smoothness of the loss functions and the scale of gradients.

Assumptions. To control the dynamics, we impose two standard conditions on the geometry of the batch objectives and the scale of pairwise gradients.

Assumption 1 (Uniform smoothness). *For every batch \mathcal{B} , the functions $w \mapsto \bar{\ell}_{\mathcal{B}}^{\text{CL}}(w)$ and $w \mapsto \bar{\ell}_{\mathcal{B}}^{\text{NSCL}}(w)$ are β -smooth with the same constant $\beta > 0$:*

$$\|\nabla \phi(w) - \nabla \phi(v)\| \leq \beta \|w - v\| \quad \text{for all } v, w \in \mathbb{R}^p, \phi \in \{\bar{\ell}_{\mathcal{B}}^{\text{CL}}, \bar{\ell}_{\mathcal{B}}^{\text{NSCL}}\}.$$

Assumption 2 (Bounded pairwise gradients). *There exists $G > 0$, independent of \mathcal{B} and t , such that for all w and all pairs (u, v) appearing in any denominator term,*

$$\|\nabla_w \text{sim}(f_w(u), f_w(v))\| \leq G.$$

We quantify drift between the coupled trajectories under shared randomness in the *nonconvex β -smooth* regime. Throughout, the only data-dependent term is $\Delta_{C,\delta}(B; \tau)$, which decreases with more classes and larger batches.

Theorem 2. *Fix $B, T \in \mathbb{N}$, $\delta \in (0, 1)$, and temperature $\tau > 0$. Suppose Assumptions 1–2 hold. Then, with probability at least $1 - \delta$,*

$$e_T \leq \frac{G}{\beta \tau} \Delta_{C,\delta}(B; \tau) \left(\exp\left(\beta \sum_{t=0}^{T-1} \eta_t\right) - 1 \right).$$

The bound scales linearly with G and $\Delta_{C,\delta}(B; \tau)$, but crucially it is amplified by the exponential factor $\exp(\beta \sum_t \eta_t)$. Unless the step sizes are aggressively annealed, this term grows rapidly with training time. Even though $\Delta_{C,\delta}(B; \tau)$ improves with C and B (e.g., for $C=1000$, $B=512$, $\delta=0.01$, we obtain $\Delta_{C,\delta}(B; \tau) \approx 0.01$ so that the reweightings of the steps differ by about one percent), the exponential accumulation can still overwhelm this small per-step gap.

In other words, parameter-space coupling guarantees only that the two runs do not drift apart too quickly in weight space. But because the weights may follow very different trajectories even

when representations remain similar, this control is too weak to yield meaningful statements about representational alignment. This motivates our next step: shifting the analysis to *similarity space*, where we can obtain bounds that remain stable throughout training and translate directly into guarantees on metrics such as CKA and RSA.

Proof idea. With high probability over batches (Cor. 3), every anchor’s denominator is dominated by negatives up to $\epsilon_{B,\delta}$ fluctuations. This keeps the (temperature- τ) softmax reweighting gap between CL and NSCL small. In particular, Lem. 7 shows that the per-batch parameter gradients differ uniformly as

$$\|\nabla \bar{\ell}_{\mathcal{B}_t}^{\text{CL}}(w) - \nabla \bar{\ell}_{\mathcal{B}_t}^{\text{NSCL}}(w)\| \leq \frac{G}{\tau} \Delta_{C,\delta}(B; \tau).$$

By β -smoothness of each batch loss, each step can expand distances by at most a factor $(1 + \beta \eta_t)$. Combining this smoothness expansion with the uniform gradient-gap bound yields the following recurrence:

$$e_{t+1} \leq (1 + \beta \eta_t) e_t + \eta_t \frac{G}{\tau} \Delta_{C,\delta}(B; \tau),$$

where the first term propagates the previous error (with amplification controlled by curvature), and the second injects the new discrepancy introduced by the CL–NSCL gap at temperature τ .

Unrolling over T steps and applying the discrete Grönwall inequality gives the exponential-type bound

$$e_T \leq \frac{G}{\beta \tau} \Delta_{C,\delta}(B; \tau) \left(\exp\left(\beta \sum_{t=0}^{T-1} \eta_t\right) - 1 \right).$$

Thus, cumulative drift scales with the reweighting gap and is amplified exponentially with the total step size; smaller τ tightens the softmax and increases the constants (via both $1/\tau$ and $e^{2/\tau}$ inside $\Delta_{C,\delta}$), so keeping $\sum_t \eta_t$ moderate is especially important.

C Why gradient descent in similarity space is a faithful surrogate

We now explain why running gradient descent directly in similarity space closely tracks the dynamics induced by gradient descent in parameter space.

When parameters move from w_t to w_{t+1} , the induced change in the similarity matrix can be approximated by a linear expansion:

$$\Sigma(w_{t+1}) - \Sigma(w_t) \approx J_t(w_{t+1} - w_t), \quad J_t := J(w_t), \quad (3)$$

where $J(w) := \partial \Sigma / \partial w$ is the Jacobian. The error in this expansion, denoted R_t , is quadratic in the step size:

$$\Sigma(w_{t+1}) - \Sigma(w_t) = J_t(w_{t+1} - w_t) + R_t. \quad (4)$$

By the chain rule, the gradient in parameter space can be written as follows:

$$\nabla_w \bar{\ell}(w_t) = J_t^\top \nabla_\Sigma \bar{\ell}(\Sigma(w_t)) = J_t^\top \hat{G}_t,$$

where $\hat{G}_t := \nabla_\Sigma \bar{\ell}(\Sigma(w_t))$. Substituting this into the update rule gives

$$\Sigma(w_{t+1}) - \Sigma(w_t) = -\eta_t P_t \hat{G}_t + R_t, \quad P_t := J_t J_t^\top \succeq 0. \quad (5)$$

Thus, parameter descent acts like similarity descent, but with a preconditioning matrix P_t , plus the remainder R_t .

Assume there exist constants $L_\Sigma, M_\Sigma > 0$ such that

$$\|J(w)\|_{2 \rightarrow 2} \leq L_\Sigma, \quad \|\Sigma(w + \Delta w) - \Sigma(w) - J(w)\Delta w\|_F \leq \frac{M_\Sigma}{2} \|\Delta w\|_2^2.$$

Then $\|P_t\|_{2 \rightarrow 2} \leq L_\Sigma^2$ and, with $\Delta w_t := -\eta_t \nabla_w \bar{\ell}(w_t)$,

$$\|R_t\|_F \leq \frac{M_\Sigma}{2} \eta_t^2 \|\nabla_w \bar{\ell}(w_t)\|_2^2 =: \frac{M_\Sigma}{2} \eta_t^2 \Xi_t. \quad (6)$$

Let $\widehat{\Sigma}_t := \Sigma(w_t)$ be the similarity trajectory induced by parameter descent. Define $\widetilde{\Sigma}_t$ as the trajectory of explicit similarity descent:

$$\widetilde{\Sigma}_{t+1} = \widetilde{\Sigma}_t - \eta_t \widetilde{G}_t, \quad \widetilde{G}_t := \nabla_{\Sigma} \bar{\ell}(\widetilde{\Sigma}_t),$$

with $\widehat{\Sigma}_0 = \widetilde{\Sigma}_0$. Let $E_t := \|\widehat{\Sigma}_t - \widetilde{\Sigma}_t\|_F$ and $C_{\Sigma} := \sup_t \|P_t - I\|_{2 \rightarrow 2} \leq L_{\Sigma}^2 + 1$. Using equation 5, adding and subtracting $-\eta_t \widehat{G}_t$, and applying the temperature- τ bounds equation 11 and equation 6, one obtains

$$E_{t+1} \leq \left(1 + \frac{\eta_t}{2\tau^2 B}\right) E_t + \eta_t C_{\Sigma} \|\widehat{G}_t\|_F + \frac{M_{\Sigma}}{2} \eta_t^2 \Xi_t. \quad (7)$$

Unrolling this recursion from $E_0 = 0$ and using $\prod_u (1 + \alpha_u) \leq \exp(\sum_u \alpha_u)$ yields

$$\|\widehat{\Sigma}_T - \widetilde{\Sigma}_T\|_F \leq \exp\left(\frac{1}{2\tau^2 B} \sum_{t=0}^{T-1} \eta_t\right) \left[C_{\Sigma} \sum_{t=0}^{T-1} \eta_t \|\widehat{G}_t\|_F + \frac{M_{\Sigma}}{2} \sum_{t=0}^{T-1} \eta_t^2 \Xi_t \right]. \quad (8)$$

By bounding $\|\widehat{G}_t\|_F$ via equation 13, namely $\|\widehat{G}_t\|_F \leq \frac{1}{\tau} \sqrt{\frac{2}{B}}$, this simplifies to

$$\|\widehat{\Sigma}_T - \widetilde{\Sigma}_T\|_F \leq \exp\left(\frac{1}{2\tau^2 B} \sum_{t=0}^{T-1} \eta_t\right) \left[\frac{\sqrt{2} C_{\Sigma}}{\tau \sqrt{B}} \sum_{t=0}^{T-1} \eta_t + \frac{M_{\Sigma}}{2} \sum_{t=0}^{T-1} \eta_t^2 \Xi_t \right]. \quad (9)$$

To summarize, the difference between the two trajectories stays small when step sizes are not too aggressive (moderate cumulative step size), the schedule is square-summable (so $\sum_t \eta_t^2$ remains finite), and the batch size B is not too small. The temperature τ modulates both curvature (via $1/\tau^2$) and gradient magnitudes (via $1/\tau$); smaller τ tightens the softmax and makes the coupling more sensitive to step size.

D Technical Tools and Proofs

D.1 Notation and basic softmax facts

Let $S = \{(x_i, y_i)\}_{i=1}^N$ be class-balanced with C classes and $N = Cn$ (each class has n points). For parameters w , let $z_i = f_w(x_i)$ and define the bounded similarity matrix

$$\Sigma(w)_{ij} := \text{sim}(z_i, z_j) \in [-1, 1].$$

At step t , draw a mini-batch $\mathcal{B}_t = \{(x_{j_s}, x'_{j_s}, y_{j_s})\}_{s=1}^B$ with replacement, using independent augmentations $x'_{j_s} \sim \alpha(x_{j_s})$. For an anchor $i \in \{j_1, \dots, j_B\}$, let D_i be its denominator index set, and let $D_i^{\text{neg}} := \{k \in D_i : y_k \neq y_i\}$ (and similarly D_i^{pos}) denote the subset restricted to negatives (e.g., in two-view SimCLR, D_i consists of all $2B$ views except the anchor itself).

Define the anchor's logit vector $s_i(w) := (\Sigma(w)_{i,k})_{k \in D_i}$ and the corresponding softmax distributions with temperature $\tau > 0$ (default 1):

$$p_i = \text{softmax}(s_i(w)/\tau), \quad q_i = \text{softmax}((s_i(w))_{D_i^{\text{neg}}} / \tau).$$

Let i' denote the positive (augmented) index for anchor i .

For contrastive learning (CL) and negatives-only supervised contrastive learning (NSCL), the per-anchor and batch losses are

$$\begin{aligned} \ell_i^{\text{CL}}(s_i) &= -\log p_{i,i'}, & \ell_i^{\text{NSCL}}(s_i) &= -\log q_{i,i'}, \\ \bar{\ell}_{\mathcal{B}_t}^{\text{CL}} &= \frac{1}{B} \sum_{i \in \{j_1, \dots, j_B\}} \ell_i^{\text{CL}}(s_i), & \bar{\ell}_{\mathcal{B}_t}^{\text{NSCL}} &= \frac{1}{B} \sum_{i \in \{j_1, \dots, j_B\}} \ell_i^{\text{NSCL}}(s_i). \end{aligned}$$

Since $\Sigma(w)_{ij} \in [-1, 1]$, each exponential term inside the softmax lies in

$$\exp(\Sigma(w)_{ij}/\tau) \in [e^{-1/\tau}, e^{1/\tau}],$$

a fact used below to control softmax mass ratios.

Lemma 1 (Anchor-block orthogonality). *Fix a step t and batch \mathcal{B}_t . For each anchor $i \in \mathcal{B}_t$, let D_i be the set of indices appearing in i 's denominator and define the per-anchor gradient $g_i \in \mathbb{R}^{\mathcal{I}_t}$ by*

$$g_i := \nabla_{s_i} \ell_i \quad \text{placed on the coordinates } \{(i, k) : k \in D_i\} \subset \mathcal{I}_t,$$

with zeros elsewhere (here \mathcal{I}_t is the set of all coordinates touched at step t). If $i \neq j$, then g_i and g_j have disjoint supports, and hence

$$\langle g_i, g_j \rangle_F = 0.$$

Consequently, for the batch gradient $G = \frac{1}{B} \sum_{i \in \mathcal{B}_t} g_i$,

$$\|G\|_F^2 = \frac{1}{B^2} \sum_{i \in \mathcal{B}_t} \|g_i\|_F^2. \quad (10)$$

Proof. By construction, g_i is supported only on coordinates $\{(i, k) : k \in D_i\}$, while g_j is supported only on $\{(j, k) : k \in D_j\}$. For $i \neq j$ these sets are disjoint, so every coordinatewise product is zero, yielding $\langle g_i, g_j \rangle_F = 0$. Expanding the square for G ,

$$\|G\|_F^2 = \left\langle \frac{1}{B} \sum_i g_i, \frac{1}{B} \sum_j g_j \right\rangle_F = \frac{1}{B^2} \sum_i \|g_i\|_F^2 + \frac{1}{B^2} \sum_{i \neq j} \langle g_i, g_j \rangle_F = \frac{1}{B^2} \sum_i \|g_i\|_F^2,$$

where the cross terms vanish by orthogonality. \square

Lemma 2 (Softmax Hessian and gradient Lipschitzness). *Fix a step t and batch \mathcal{B}_t . Let \mathcal{I}_t be the set of coordinates (i, k) that appear in any anchor's denominator at step t , and view $\bar{\ell}_{\mathcal{B}_t}$ (either CL or NSCL) as a function of the restricted similarity entries $\Sigma \in \mathbb{R}^{\mathcal{I}_t}$. For each anchor i , write $s_i = \{\Sigma(i, k) : (i, k) \in \mathcal{I}_t\}$ and $p_i = \text{softmax}(s_i/\tau)$. Then:*

$$\nabla_{s_i}^2 \ell_i(s_i) = \frac{1}{\tau^2} J(s_i), \quad J(s_i) := \text{Diag}(p_i) - p_i p_i^\top, \quad \|\nabla^2 \bar{\ell}_{\mathcal{B}_t}(\Sigma)\|_{2 \rightarrow 2} \leq \frac{1}{2\tau^2 B}.$$

Consequently, for all $\Sigma, \tilde{\Sigma} \in \mathbb{R}^{\mathcal{I}_t}$,

$$\|\nabla_{\Sigma} \bar{\ell}_{\mathcal{B}_t}(\Sigma) - \nabla_{\Sigma} \bar{\ell}_{\mathcal{B}_t}(\tilde{\Sigma})\|_F \leq \frac{1}{2\tau^2 B} \|\Sigma - \tilde{\Sigma}\|_F. \quad (11)$$

Proof. With temperature $\tau > 0$, for an anchor i we have $p_i = \text{softmax}(s_i/\tau)$ and

$$\nabla_{s_i} \ell_i(s_i) = \frac{1}{\tau} (p_i - e_{i'}) \implies \nabla_{s_i}^2 \ell_i(s_i) = \frac{1}{\tau^2} \nabla_{s_i} p_i = \frac{1}{\tau^2} J(s_i),$$

where $J(s_i) := \text{Diag}(p_i) - p_i p_i^\top$. Bound $\|J(s_i)\|_{2 \rightarrow 2}$ via the infinity norm:

$$\begin{aligned} \|J(s_i)\|_{2 \rightarrow 2} &\leq \|J(s_i)\|_\infty \\ &= \max_r \sum_\ell |J_{r\ell}| \\ &= \max_r \left(p_{i,r} (1 - p_{i,r}) + \sum_{\ell \neq r} p_{i,r} p_{i,\ell} \right) \\ &= \max_r 2p_{i,r}(1 - p_{i,r}) \leq \frac{1}{2}, \end{aligned}$$

since $x(1-x) \leq 1/4$ for $x \in [0, 1]$.

The batch loss is an average over anchors, so its Hessian is block-diagonal across anchors with a prefactor $1/B$:

$$\nabla^2 \bar{\ell}_{\mathcal{B}_t}(\Sigma) = \frac{1}{B} \text{blkdiag} \left(\frac{1}{\tau^2} J(s_i) \right)_{i \in \mathcal{B}_t} = \frac{1}{\tau^2 B} \text{blkdiag} (J(s_i))_{i \in \mathcal{B}_t}.$$

Hence

$$\|\nabla^2 \bar{\ell}_{\mathcal{B}_t}(\Sigma)\|_{2 \rightarrow 2} = \frac{1}{\tau^2 B} \max_i \|J(s_i)\|_{2 \rightarrow 2} \leq \frac{1}{2\tau^2 B}.$$

By the mean-value (integral) form for vector fields,

$$\nabla_{\Sigma} \bar{\ell}_{\mathcal{B}_t}(\Sigma) - \nabla_{\Sigma} \bar{\ell}_{\mathcal{B}_t}(\tilde{\Sigma}) = \int_0^1 \nabla^2 \bar{\ell}_{\mathcal{B}_t}(\tilde{\Sigma} + \theta(\Sigma - \tilde{\Sigma})) [\Sigma - \tilde{\Sigma}] d\theta,$$

and therefore

$$\|\nabla_{\Sigma} \bar{\ell}_{\mathcal{B}_t}(\Sigma) - \nabla_{\Sigma} \bar{\ell}_{\mathcal{B}_t}(\tilde{\Sigma})\|_F \leq \sup_{\theta \in [0, 1]} \|\nabla^2 \bar{\ell}_{\mathcal{B}_t}(\Sigma_\theta)\|_{2 \rightarrow 2} \|\Sigma - \tilde{\Sigma}\|_F \leq \frac{1}{2\tau^2 B} \|\Sigma - \tilde{\Sigma}\|_F,$$

as claimed. \square

Lemma 3 (Per-anchor gradient norm and batch average). *For an anchor i , let s_i be the vector of logits in its denominator and $p_i = \text{softmax}(s_i/\tau)$. Let i' denote the (unique) positive index (for NSCL, if i' is not in the denominator, set $p_{i,i'} := 0$ in the display below). Then*

$$\|\nabla_{s_i} \ell_i\|_2^2 = \frac{1}{\tau^2} \left[(1 - p_{i,i'})^2 + \sum_{k \neq i'} p_{i,k}^2 \right] \leq \frac{2}{\tau^2}, \quad (12)$$

hence $\|\nabla_{s_i} \ell_i\|_2 \leq \sqrt{2}/\tau$. Moreover, by block orthogonality across anchors,

$$\left\| \frac{1}{B} \sum_{i \in \mathcal{B}_t} \nabla_{s_i} \ell_i \right\|_F^2 = \frac{1}{B^2} \sum_{i \in \mathcal{B}_t} \|\nabla_{s_i} \ell_i\|_2^2 \leq \frac{2}{\tau^2 B} \implies \left\| \frac{1}{B} \sum_{i \in \mathcal{B}_t} \nabla_{s_i} \ell_i \right\|_F \leq \frac{1}{\tau} \sqrt{\frac{2}{B}}. \quad (13)$$

Proof. For CL, the loss is $-\log p_{i,i'}$ with $p_i = \text{softmax}(s_i/\tau)$. By the standard softmax–cross-entropy derivative with temperature,

$$\nabla_{s_i} \ell_i = \frac{1}{\tau} (p_i - e_{i'}),$$

so

$$\|\nabla_{s_i} \ell_i\|_2^2 = \frac{1}{\tau^2} \left[(1 - p_{i,i'})^2 + \sum_{k \neq i'} p_{i,k}^2 \right] \leq \frac{1}{\tau^2} \left[(1 - p_{i,i'})^2 + \left(\sum_{k \neq i'} p_{i,k} \right)^2 \right] = \frac{2}{\tau^2} (1 - p_{i,i'})^2 \leq \frac{2}{\tau^2},$$

since p_i is a probability vector and $\sum_{k \neq i'} p_{i,k} = 1 - p_{i,i'}$.

For NSCL, two cases. If $i' \in D_i$, the same computation applies (the target index is present), hence the same bound holds. If $i' \notin D_i$ (negatives-only denominator), then the loss is $-\log q_{i,i'}$ with $q_i = \text{softmax}((s_i)_{D_i^-}/\tau)$ supported only on D_i^- , and

$$\nabla_{s_i} \ell_i = \frac{1}{\tau} q_i \quad \text{on } D_i^- \quad (\text{and 0 on } D_i^+),$$

so

$$\|\nabla_{s_i} \ell_i\|_2^2 = \frac{1}{\tau^2} \sum_{j \in D_i^-} q_{i,j}^2 \leq \frac{1}{\tau^2} \left(\sum_{j \in D_i^-} q_{i,j} \right)^2 = \frac{1}{\tau^2} \leq \frac{2}{\tau^2}.$$

Thus in all cases $\|\nabla_{s_i} \ell_i\|_2 \leq \sqrt{2}/\tau$, establishing equation 12.

For the batch bound equation 13, gradients from different anchors have disjoint supports over coordinates $\{(i, k) : k \in D_i\}$, so they are orthogonal in Frobenius inner product (Lem. 1). Therefore,

$$\left\| \frac{1}{B} \sum_{i \in \mathcal{B}_t} \nabla_{s_i} \ell_i \right\|_F^2 = \frac{1}{B^2} \sum_{i \in \mathcal{B}_t} \|\nabla_{s_i} \ell_i\|_2^2 \leq \frac{1}{B^2} \cdot B \cdot \frac{2}{\tau^2} = \frac{2}{\tau^2 B},$$

which also implies $\left\| \frac{1}{B} \sum_{i \in \mathcal{B}_t} \nabla_{s_i} \ell_i \right\|_F \leq \frac{1}{\tau} \sqrt{2/B}$. \square

Lemma 4 (Bounded logits imply bounded softmax masses). *Fix a step t and an anchor i . Suppose all active logits satisfy $\Sigma(i, k) \in [-1, 1]$. For any index subset S in the anchor's denominator, define*

$$Z_S := \sum_{k \in S} \exp(\Sigma(i, k)/\tau) \quad \text{with temperature } \tau > 0.$$

Then

$$|S| e^{-1/\tau} \leq Z_S \leq |S| e^{1/\tau}.$$

In particular, if S_{pos} and S_{neg} are the positive and negative index sets with sizes n_{pos} and n_{neg} , and $Z_{\text{pos}} := Z_{S_{\text{pos}}}$, $Z_{\text{neg}} := Z_{S_{\text{neg}}}$, then

$$n_{\text{pos}} e^{-1/\tau} \leq Z_{\text{pos}} \leq n_{\text{pos}} e^{1/\tau}, \quad n_{\text{neg}} e^{-1/\tau} \leq Z_{\text{neg}} \leq n_{\text{neg}} e^{1/\tau},$$

and hence

$$\frac{Z_{\text{pos}}}{Z_{\text{neg}}} \leq e^{2/\tau} \frac{n_{\text{pos}}}{n_{\text{neg}}} \quad \text{and} \quad \frac{Z_{\text{pos}}}{Z_{\text{neg}}} \geq e^{-2/\tau} \frac{n_{\text{pos}}}{n_{\text{neg}}}.$$

Proof. Since $\Sigma(i, k) \in [-1, 1]$, we have $\exp(\Sigma(i, k)/\tau) \in [e^{-1/\tau}, e^{1/\tau}]$ for every active k . Summing over $k \in S$ yields $|S| e^{-1/\tau} \leq Z_S \leq |S| e^{1/\tau}$. Apply this with $S = S_{\text{pos}}$ and $S = S_{\text{neg}}$ and take ratios to obtain the stated bounds. \square

D.2 High-probability batch composition

Fix $T, B \in \mathbb{N}$ and $\epsilon > 0$. For step t and anchor $i \in \mathcal{B}_t$, let $Y_{t,s}^{(i)} = \mathbf{1}\{y_{j_s} \neq y_i\}$ for $s = 1, \dots, B$.

Lemma 5 (Batch-composition event). *For a class-balanced population with C classes, the $Y_{t,s}^{(i)}$ are i.i.d. Bernoulli with mean $1 - \frac{1}{C}$. For any $\epsilon > 0$,*

$$\mathbb{P} \left[\exists(t, i) : \frac{1}{B} \sum_{s=1}^B Y_{t,s}^{(i)} < 1 - \frac{1}{C} - \epsilon \right] \leq TB e^{-2B\epsilon^2}.$$

Equivalently, with probability $\geq 1 - TB e^{-2B\epsilon^2}$, every anchor sees at least $B(1 - \frac{1}{C} - \epsilon)$ negatives.

Proof. Fix any step t and anchor i . Because batches are drawn with replacement from a class-balanced population with C classes, for each position $s \in \{1, \dots, B\}$ the indicator $Y_{t,s}^{(i)} = \mathbf{1}\{y_{j_s} \neq y_i\}$ is Bernoulli with mean $\mathbb{E}[Y_{t,s}^{(i)}] = 1 - \frac{1}{C}$, and $\{Y_{t,s}^{(i)}\}_{s=1}^B$ are i.i.d. across s . By Hoeffding's inequality, for any $\epsilon > 0$,

$$\mathbb{P} \left[\frac{1}{B} \sum_{s=1}^B Y_{t,s}^{(i)} < 1 - \frac{1}{C} - \epsilon \right] = \mathbb{P} \left[\frac{1}{B} \sum_{s=1}^B (Y_{t,s}^{(i)} - \mathbb{E}Y_{t,s}^{(i)}) < -\epsilon \right] \leq \exp(-2B\epsilon^2).$$

There are at most TB anchor-step pairs (t, i) over $t = 0, \dots, T-1$ and $i \in \mathcal{B}_t$. A union bound gives

$$\mathbb{P} \left[\exists(t, i) : \frac{1}{B} \sum_{s=1}^B Y_{t,s}^{(i)} < 1 - \frac{1}{C} - \epsilon \right] \leq TB e^{-2B\epsilon^2}.$$

Equivalently, with probability at least $1 - TB e^{-2B\epsilon^2}$, every anchor in every step has at least $B(1 - \frac{1}{C} - \epsilon)$ negatives in its denominator. \square

Corollary 3. *For $\delta \in (0, 1)$, set $\epsilon_{B,\delta} := \sqrt{\frac{1}{2B} \log(\frac{TB}{\delta})}$. With probability $\geq 1 - \delta$, every anchor i has at least $B(1 - \frac{1}{C} - \epsilon_{B,\delta})$ negatives and at most $B(\frac{1}{C} + \epsilon_{B,\delta})$ positives in its denominator. Using bounded logits, the ratio of total positive to negative softmax mass (at temperature $\tau > 0$) satisfies, for all anchors and steps,*

$$\frac{Z_i^{\text{pos}}}{Z_i^{\text{neg}}} \leq \frac{e^{2/\tau} (\frac{1}{C} + \epsilon_{B,\delta})}{1 - \frac{1}{C} - \epsilon_{B,\delta}} = \frac{1}{2} \Delta_{C,\delta}(B; \tau), \quad (14)$$

where

$$\Delta_{C,\delta}(B; \tau) := \frac{2 e^{2/\tau} (\frac{1}{C} + \epsilon_{B,\delta})}{1 - \frac{1}{C} - \epsilon_{B,\delta}}.$$

Proof. Set $\epsilon = \epsilon_{B,\delta} := \sqrt{\frac{1}{2B} \log(\frac{TB}{\delta})}$ and $\Delta_{C,\delta}(B; \tau) := \frac{2 e^{2/\tau} (\frac{1}{C} + \epsilon_{B,\delta})}{1 - \frac{1}{C} - \epsilon_{B,\delta}}$. Apply Lem. 5 with this ϵ : with probability at least $1 - \delta$, for every step t and every anchor i ,

$$|D_i^{\text{neg}}| \geq B \left(1 - \frac{1}{C} - \epsilon_{B,\delta} \right), \quad |D_i^{\text{pos}}| \leq B \left(\frac{1}{C} + \epsilon_{B,\delta} \right).$$

In two-view SimCLR, each sampled point contributes two denominator entries, so the denominator contains at least $2|D_i^{\text{neg}}|$ negative entries and at most $2|D_i^{\text{pos}}|$ positive entries; the factor 2 cancels in the ratio below.

Because similarities are bounded in $[-1, 1]$, each logit lies in $[-1, 1]$ and hence each exponential term at temperature τ lies in $[e^{-1/\tau}, e^{1/\tau}]$. Therefore, for any anchor and step,

$$Z_i^{\text{pos}} \leq e^{1/\tau} \cdot (2|D_i^{\text{pos}}|), \quad Z_i^{\text{neg}} \geq e^{-1/\tau} \cdot (2|D_i^{\text{neg}}|),$$

and thus

$$\frac{Z_i^{\text{pos}}}{Z_i^{\text{neg}}} \leq e^{2/\tau} \frac{|D_i^{\text{pos}}|}{|D_i^{\text{neg}}|} \leq \frac{e^{2/\tau} \left(\frac{1}{C} + \epsilon_{B,\delta} \right)}{1 - \frac{1}{C} - \epsilon_{B,\delta}} = \frac{1}{2} \Delta_{C,\delta}(B; \tau).$$

The bound is meaningful whenever $\epsilon_{B,\delta} < 1 - \frac{1}{C}$ so that the denominator is positive. This proves the corollary. \square

Lemma 6 (Per-anchor reweighting gap). *On the event of Cor. 3, let p be the CL softmax (temperature $\tau > 0$) over an anchor's full denominator, and q the NSCL softmax (same τ) that removes same-class entries and renormalizes over negatives. Then*

$$\|p - q\|_1 \leq \Delta_{C,\delta}(B; \tau), \quad \|p - q\|_2 \leq \|p - q\|_1 \leq \Delta_{C,\delta}(B; \tau).$$

Proof. Fix an anchor i and let $D_i^{\text{pos}}, D_i^{\text{neg}}$ be its positive and negative index sets in the CL denominator. Write $s_k := \Sigma(i, k)$ and define

$$Z_i^{\text{pos}} := \sum_{k \in D_i^{\text{pos}}} \exp(s_k / \tau), \quad Z_i^{\text{neg}} := \sum_{j \in D_i^{\text{neg}}} \exp(s_j / \tau), \quad \alpha := \frac{Z_i^{\text{pos}}}{Z_i^{\text{pos}} + Z_i^{\text{neg}}}.$$

Let p be the CL softmax on $D_i^{\text{pos}} \cup D_i^{\text{neg}}$ and let q be the NSCL softmax that zeros positive entries and renormalizes on negatives: $q(k) = 0$ for $k \in D_i^{\text{pos}}$ and $q(j) = p(j)/(1 - \alpha)$ for $j \in D_i^{\text{neg}}$. Then

$$\|p - q\|_1 = \sum_{k \in D_i^{\text{pos}}} p_k + \sum_{j \in D_i^{\text{neg}}} \left| p_j - \frac{p_j}{1 - \alpha} \right| = \alpha + (1 - \alpha) \frac{\alpha}{1 - \alpha} = 2\alpha \leq \frac{2Z_i^{\text{pos}}}{Z_i^{\text{neg}}}.$$

On the high-probability event of Cor. 3, since $s \in [-1, 1] \Rightarrow \exp(s/\tau) \in [e^{-1/\tau}, e^{1/\tau}]$,

$$Z_i^{\text{pos}} \leq e^{1/\tau} |D_i^{\text{pos}}|, \quad Z_i^{\text{neg}} \geq e^{-1/\tau} |D_i^{\text{neg}}|.$$

Moreover,

$$|D_i^{\text{pos}}| \leq 2B \left(\frac{1}{C} + \epsilon_{B,\delta} \right), \quad |D_i^{\text{neg}}| \geq 2B \left(1 - \frac{1}{C} - \epsilon_{B,\delta} \right),$$

(each sampled point contributes two keys, so the factor 2 cancels in the ratio). Hence

$$\frac{2Z_i^{\text{pos}}}{Z_i^{\text{neg}}} \leq 2e^{2/\tau} \frac{|D_i^{\text{pos}}|}{|D_i^{\text{neg}}|} \leq \frac{2e^{2/\tau} \left(\frac{1}{C} + \epsilon_{B,\delta} \right)}{1 - \frac{1}{C} - \epsilon_{B,\delta}} =: \Delta_{C,\delta}(B; \tau).$$

Therefore $\|p - q\|_1 \leq \Delta_{C,\delta}(B; \tau)$. Finally, $\|p - q\|_2 \leq \|p - q\|_1$ yields the second claim. \square

D.3 Parameter-space coupling: supporting lemmas and proofs

Lemma 7 (Per-batch parameter-gradient gap). *On the event of Cor. 3, for any step t and any w ,*

$$\|\nabla_w \bar{\ell}_{\mathcal{B}_t}^{\text{CL}}(w) - \nabla_w \bar{\ell}_{\mathcal{B}_t}^{\text{NSCL}}(w)\| \leq \frac{G}{\tau} \Delta_{C,\delta}(B; \tau).$$

Proof. Fix t and w . For an anchor $i \in \mathcal{B}_t$, let D_i be its denominator index set, split as $D_i = \text{pos}_i \cup \text{neg}_i$, where pos_i collects all same-class indices (including the designated positive i') and neg_i the rest. Write the logits $s_{ik} = \Sigma(i, k)$, the CL softmax $p_{ik} = \exp(s_{ik}/\tau) / \sum_{\ell \in D_i} \exp(s_{i\ell}/\tau)$, and the NSCL softmax over negatives $q_{ij} = p_{ij}/(1 - \alpha_i)$ for $j \in \text{neg}_i$, with $q_k = 0$ for $k \in \text{pos}_i$, where $\alpha_i := \sum_{k \in \text{pos}_i} p_{ik}$. Define $v_{ik} := \nabla_w s_{ik} = \nabla_w \text{sim}(f_w(x_i), f_w(x_k))$; by Assumption 2, $\|v_{ik}\| \leq G$ for all (i, k) .

For the per-anchor losses,

$$\nabla_w \ell_{i,\mathcal{B}_t}^{\text{CL}} = \frac{1}{\tau} \left(\sum_{k \in D_i} p_{ik} v_{ik} - v_{ii'} \right), \quad \nabla_w \ell_{i,\mathcal{B}_t}^{\text{NSCL}} = \frac{1}{\tau} \left(\sum_{j \in \text{neg}_i} q_{ij} v_{ij} - v_{ii'} \right).$$

Hence the per-anchor gradient difference is

$$\Delta g_i := \nabla_w \ell_{i,\mathcal{B}_t}^{\text{CL}} - \nabla_w \ell_{i,\mathcal{B}_t}^{\text{NSCL}} = \frac{1}{\tau} \left(\underbrace{\sum_{k \in \text{pos}_i} p_{ik} v_{ik}}_{(\text{A})} + \underbrace{\sum_{j \in \text{neg}_i} (p_{ij} - q_{ij}) v_{ij}}_{(\text{B})} \right).$$

By the triangle inequality and $\|v_{ik}\| \leq G$,

$$\|\Delta g_i\| \leq \frac{G}{\tau} \left(\sum_{k \in \text{pos}_i} p_{ik} + \sum_{j \in \text{neg}_i} |p_{ij} - q_{ij}| \right).$$

Since $q_{ij} = p_{ij}/(1 - \alpha_i)$ for $j \in \text{neg}_i$,

$$\sum_{j \in \text{neg}_i} |p_{ij} - q_{ij}| = \sum_{j \in \text{neg}_i} p_{ij} \frac{\alpha_i}{1 - \alpha_i} = \alpha_i.$$

Therefore $\|\Delta g_i\| \leq \frac{G}{\tau} (\alpha_i + \alpha_i) = \frac{2G}{\tau} \alpha_i$. Writing $r_i := \frac{Z_{\text{pos}}}{Z_{\text{neg}}}$ with $Z_{\text{pos}} = \sum_{k \in \text{pos}_i} \exp(s_{ik}/\tau)$, $Z_{\text{neg}} = \sum_{j \in \text{neg}_i} \exp(s_{ij}/\tau)$, we have $\alpha_i = \frac{r_i}{1+r_i}$, hence $2\alpha_i = \frac{2r_i}{1+r_i} \leq 2r_i$, so

$$\|\Delta g_i\| \leq \frac{2G}{\tau} \frac{Z_{\text{pos}}}{Z_{\text{neg}}}.$$

On the high-probability event of Cor. 3, for every anchor

$$\frac{Z_{\text{pos}}}{Z_{\text{neg}}} \leq \frac{e^{2/\tau} \left(\frac{1}{C} + \epsilon_{B,\delta} \right)}{1 - \frac{1}{C} - \epsilon_{B,\delta}} = \frac{1}{2} \Delta_{C,\delta}(B; \tau),$$

so $\|\Delta g_i\| \leq \frac{G}{\tau} \Delta_{C,\delta}(B; \tau)$ for all anchors i .

Finally, the batch gradients are averages over anchors:

$$\nabla \bar{\ell}_{\mathcal{B}_t}^{\text{CL}} - \nabla \bar{\ell}_{\mathcal{B}_t}^{\text{NSCL}} = \frac{1}{B} \sum_{i \in \mathcal{B}_t} \Delta g_i,$$

hence

$$\|\nabla \bar{\ell}_{\mathcal{B}_t}^{\text{CL}} - \nabla \bar{\ell}_{\mathcal{B}_t}^{\text{NSCL}}\| \leq \frac{1}{B} \sum_{i \in \mathcal{B}_t} \|\Delta g_i\| \leq \frac{1}{B} \sum_{i \in \mathcal{B}_t} \frac{G}{\tau} \Delta_{C,\delta}(B; \tau) = \frac{G}{\tau} \Delta_{C,\delta}(B; \tau).$$

□

Theorem 2. Fix $B, T \in \mathbb{N}$, $\delta \in (0, 1)$, and temperature $\tau > 0$. Suppose Assumptions 1–2 hold. Then, with probability at least $1 - \delta$,

$$e_T \leq \frac{G}{\beta \tau} \Delta_{C,\delta}(B; \tau) \left(\exp \left(\beta \sum_{t=0}^{T-1} \eta_t \right) - 1 \right).$$

Proof. Let $\Phi_t^{\text{CL}}(w) := \bar{\ell}_{\mathcal{B}_t}^{\text{CL}}(w)$ and $\Phi_t^{\text{NSCL}}(w) := \bar{\ell}_{\mathcal{B}_t}^{\text{NSCL}}(w)$. Assume each Φ_t^{CL} is β -smooth. Set $e_t := \|w_t^{\text{CL}} - w_t^{\text{NSCL}}\|$.

Write

$$\begin{aligned} e_{t+1} &= \|w_{t+1}^{\text{CL}} - w_{t+1}^{\text{NSCL}}\| = \|T_t(w_t^{\text{CL}}) - (w_t^{\text{NSCL}} - \eta_t \nabla \Phi_t^{\text{NSCL}}(w_t^{\text{NSCL}}))\| \\ &\leq \underbrace{\|T_t(w_t^{\text{CL}}) - T_t(w_t^{\text{NSCL}})\|}_{(\text{I})} + \eta_t \underbrace{\|\nabla \Phi_t^{\text{CL}}(w_t^{\text{NSCL}}) - \nabla \Phi_t^{\text{NSCL}}(w_t^{\text{NSCL}})\|}_{(\text{II})}. \end{aligned}$$

Bounding (I). Using the integral Hessian representation,

$$\nabla \Phi_t^{\text{CL}}(u) - \nabla \Phi_t^{\text{CL}}(v) = H_t(v, u)(u - v), \quad H_t(v, u) := \int_0^1 \nabla^2 \Phi_t^{\text{CL}}(v + \tau(u - v)) d\tau,$$

and β -smoothness gives $\|H_t(v, u)\|_{2 \rightarrow 2} \leq \beta$. Hence

$$\begin{aligned} \|T_t(u) - T_t(v)\| &= \|(I - \eta_t H_t(v, u))(u - v)\| \\ &\leq \|I - \eta_t H_t(v, u)\|_{2 \rightarrow 2} \|u - v\| \\ &\leq (1 + \eta_t \beta) \|u - v\|. \end{aligned}$$

Thus, $(I) \leq (1 + \eta_t \beta) e_t$.

Bounding (II). On the high-probability event of Cor. 3, Lem. 7 yields

$$(II) \leq \frac{G}{\tau} \Delta_{C,\delta}(B; \tau).$$

Combining the bounds,

$$e_{t+1} \leq (1 + \eta_t \beta) e_t + \eta_t \frac{G}{\tau} \Delta_{C,\delta}(B; \tau). \quad (15)$$

Iterating equation 15 from $e_0 = 0$ gives

$$e_T \leq \sum_{t=0}^{T-1} \eta_t \frac{G}{\tau} \Delta_{C,\delta}(B; \tau) \prod_{s=t+1}^{T-1} (1 + \eta_s \beta) \leq \frac{G}{\tau} \Delta_{C,\delta}(B; \tau) \sum_{t=0}^{T-1} \eta_t \exp\left(\beta \sum_{s=t+1}^{T-1} \eta_s\right),$$

where we used $1 + x \leq e^x$. Let $S_k := \sum_{s=k}^{T-1} \eta_s$ so that $S_t = \eta_t + S_{t+1}$. Then for each t ,

$$\eta_t \exp(\beta S_{t+1}) \leq \frac{1}{\beta} (\exp(\beta S_t) - \exp(\beta S_{t+1})),$$

since $e^{\beta \eta_t} - 1 \geq \beta \eta_t$. Summing over $t = 0, \dots, T-1$ telescopes to

$$e_T \leq \frac{G}{\beta \tau} \Delta_{C,\delta}(B; \tau) \left(\exp\left(\beta \sum_{t=0}^{T-1} \eta_t\right) - 1 \right).$$

This holds with probability at least $1 - \delta$ (by Cor. 3). \square

D.4 Similarity-space analysis and coupling

Lemma 8 (Per-step gradient gap in similarity space). *On the event of Cor. 3, for any step t ,*

$$\|G_t^{\text{CL}}(\Sigma_t^{\text{CL}}) - G_t^{\text{NSCL}}(\Sigma_t^{\text{NSCL}})\|_F \leq \underbrace{\frac{1}{\tau} \cdot \frac{\Delta_{C,\delta}(B; \tau)}{\sqrt{B}}}_{\text{reweighting (block-orth.)}} + \underbrace{\frac{1}{2\tau^2 B} \|\Sigma_t^{\text{CL}} - \Sigma_t^{\text{NSCL}}\|_F}_{\text{Lipschitz in } \Sigma}.$$

Proof. Add and subtract $G_t^{\text{NSCL}}(\Sigma_t^{\text{CL}})$ and apply the triangle inequality:

$$\begin{aligned} & \|G_t^{\text{CL}}(\Sigma_t^{\text{CL}}) - G_t^{\text{NSCL}}(\Sigma_t^{\text{NSCL}})\|_F \\ & \leq \underbrace{\|G_t^{\text{CL}}(\Sigma_t^{\text{CL}}) - G_t^{\text{NSCL}}(\Sigma_t^{\text{CL}})\|_F}_{(A)} + \underbrace{\|G_t^{\text{NSCL}}(\Sigma_t^{\text{CL}}) - G_t^{\text{NSCL}}(\Sigma_t^{\text{NSCL}})\|_F}_{(B)}. \end{aligned} \quad (16)$$

Term (B): Lipschitz in Σ . By the temperature- τ softmax–Hessian bound equation 11,

$$(B) \leq \frac{1}{2\tau^2 B} \|\Sigma_t^{\text{CL}} - \Sigma_t^{\text{NSCL}}\|_F.$$

Term (A): reweighting gap at fixed Σ_t^{CL} . Decompose the batch gradient into anchor blocks:

$$G_t^\circ(\Sigma) = \frac{1}{B} \sum_{i \in \mathcal{B}_t} g_{t,i}^\circ(\Sigma), \quad \circ \in \{\text{CL}, \text{NSCL}\},$$

where each $g_{t,i}^\circ$ has support only on the coordinates of anchor i . For anchor i , with temperature τ , $g_{t,i}^{\text{CL}}(\Sigma_t^{\text{CL}}) = (1/\tau)(p_i - e_{i'})$, $g_{t,i}^{\text{NSCL}}(\Sigma_t^{\text{CL}}) = (1/\tau)(q_i - e_{i'})$, so $g_{t,i}^{\text{CL}}(\Sigma_t^{\text{CL}}) - g_{t,i}^{\text{NSCL}}(\Sigma_t^{\text{CL}}) = (1/\tau)(p_i - q_i)$ on that block. By block orthogonality (Lem. 1),

$$(A) = \frac{1}{B} \left\| \sum_{i \in \mathcal{B}_t} \frac{1}{\tau} (p_i - q_i) \right\|_F = \frac{1}{\tau B} \sqrt{\sum_{i \in \mathcal{B}_t} \|p_i - q_i\|_2^2}.$$

On the event of Cor. 3, Lem. 6 gives $\|p_i - q_i\|_2 \leq \Delta_{C,\delta}(B; \tau)$ for every anchor, hence

$$(A) \leq \frac{1}{\tau B} \sqrt{B \Delta_{C,\delta}(B; \tau)^2} = \frac{1}{\tau} \cdot \frac{\Delta_{C,\delta}(B; \tau)}{\sqrt{B}}.$$

Combining the bounds on (A) and (B) yields the claim. \square

Theorem 1 (Similarity-space coupling). *Fix $B, T \in \mathbb{N}$, $\delta \in (0, 1)$, and temperature $\tau > 0$. Consider the coupled similarity-descent recursions equation 1 for CL and NSCL with shared initialization and shared mini-batches/augmentations. Then, with probability at least $1 - \delta$ over the draws of the mini-batches and augmentations, for any stepsizes $(\eta_t)_{t=0}^{T-1}$,*

$$\|\Sigma_T^{\text{CL}} - \Sigma_T^{\text{NSCL}}\|_F \leq \exp\left(\frac{1}{2\tau^2 B} \sum_{t=0}^{T-1} \eta_t\right) \frac{1}{\tau\sqrt{B}} \left(\sum_{t=0}^{T-1} \eta_t\right) \Delta_{C,\delta}(B; \tau). \quad (2)$$

Proof. Condition on the event of Cor. 3 (which holds with probability at least $1 - \delta$). Let $D_t := \|\Sigma_t^{\text{CL}} - \Sigma_t^{\text{NSCL}}\|_F$. From the coupled updates equation 1,

$$\Sigma_{t+1}^{\text{CL}} - \Sigma_{t+1}^{\text{NSCL}} = (\Sigma_t^{\text{CL}} - \Sigma_t^{\text{NSCL}}) - \eta_t (G_t^{\text{CL}}(\Sigma_t^{\text{CL}}) - G_t^{\text{NSCL}}(\Sigma_t^{\text{NSCL}})),$$

hence

$$D_{t+1} \leq D_t + \eta_t \|G_t^{\text{CL}}(\Sigma_t^{\text{CL}}) - G_t^{\text{NSCL}}(\Sigma_t^{\text{NSCL}})\|_F.$$

Add and subtract $G_t^{\text{NSCL}}(\Sigma_t^{\text{CL}})$ and apply Lem. 8 (reweighting gap + Lipschitz with temperature τ):

$$\|G_t^{\text{CL}}(\Sigma_t^{\text{CL}}) - G_t^{\text{NSCL}}(\Sigma_t^{\text{NSCL}})\|_F \leq \frac{1}{\tau} \cdot \frac{\Delta_{C,\delta}(B; \tau)}{\sqrt{B}} + \frac{1}{2\tau^2 B} D_t.$$

Therefore,

$$D_{t+1} \leq \left(1 + \frac{\eta_t}{2\tau^2 B}\right) D_t + \eta_t \frac{1}{\tau} \cdot \frac{\Delta_{C,\delta}(B; \tau)}{\sqrt{B}}.$$

Let $\alpha_t := \frac{\eta_t}{2\tau^2 B}$ and $\gamma_t := \eta_t \frac{\Delta_{C,\delta}(B; \tau)}{\tau\sqrt{B}}$. With $D_0 = 0$ (shared initialization), the discrete Grönwall/product form gives

$$D_T \leq \sum_{s=0}^{T-1} \gamma_s \prod_{u=s+1}^{T-1} (1 + \alpha_u) \leq \exp\left(\sum_{u=0}^{T-1} \alpha_u\right) \sum_{s=0}^{T-1} \gamma_s,$$

using $\prod_u (1 + \alpha_u) \leq \exp(\sum_u \alpha_u)$. Substituting α_t, γ_t yields

$$D_T \leq \exp\left(\frac{1}{2\tau^2 B} \sum_{t=0}^{T-1} \eta_t\right) \frac{1}{\tau\sqrt{B}} \left(\sum_{t=0}^{T-1} \eta_t\right) \Delta_{C,\delta}(B; \tau),$$

as desired. \square

Consequences for CKA and RSA.

Corollary 1 (CKA lower bound). *In the setting of Thm. 1. Assume $\|K_T^{\text{CL}}\|_F > 0$. With probability at least $1 - \delta$,*

$$\text{CKA}_T \geq \frac{1 - \rho_T}{1 + \rho_T}, \quad \rho_T \leq \frac{\exp\left(\frac{1}{2\tau^2 B} \sum_{t=0}^{T-1} \eta_t\right) \frac{1}{\tau\sqrt{B}} \left(\sum_{t=0}^{T-1} \eta_t\right) \Delta_{C,\delta}(B; \tau)}{\|K_T^{\text{CL}}\|_F}.$$

Proof. Let $A_T := \|K_T^{\text{CL}}\|_F > 0$ and $\Delta_{K,T} := \|K_T^{\text{CL}} - K_T^{\text{NSCL}}\|_F$, where all norms are Frobenius. Then

$$\begin{aligned} \langle K_T^{\text{CL}}, K_T^{\text{NSCL}} \rangle &= \langle K_T^{\text{CL}}, K_T^{\text{CL}} + (K_T^{\text{NSCL}} - K_T^{\text{CL}}) \rangle \\ &= \|K_T^{\text{CL}}\|_F^2 + \langle K_T^{\text{CL}}, K_T^{\text{NSCL}} - K_T^{\text{CL}} \rangle \geq A_T^2 - A_T \Delta_{K,T}, \end{aligned} \quad (17)$$

by Cauchy–Schwarz. By the triangle inequality, $\|K_T^{\text{NSCL}}\|_F \leq A_T + \Delta_{K,T}$. Hence

$$\text{CKA}_T = \frac{\langle K_T^{\text{CL}}, K_T^{\text{NSCL}} \rangle}{\|K_T^{\text{CL}}\|_F \|K_T^{\text{NSCL}}\|_F} \geq \frac{A_T^2 - A_T \Delta_{K,T}}{A_T(A_T + \Delta_{K,T})} = \frac{1 - \Delta_{K,T}/A_T}{1 + \Delta_{K,T}/A_T}.$$

Next, $K_T^\circ = H \Sigma_T^\circ H$ with the centering projector $H = I - \frac{1}{N} \mathbf{1} \mathbf{1}^\top$, so $\Delta_{K,T} = \|H(\Sigma_T^{\text{CL}} - \Sigma_T^{\text{NSCL}})H\|_F \leq \|\Sigma_T^{\text{CL}} - \Sigma_T^{\text{NSCL}}\|_F$ because $\|H\|_{2 \rightarrow 2} = 1$. By Thm. 1, with probability at least $1 - \delta$,

$$\|\Sigma_T^{\text{CL}} - \Sigma_T^{\text{NSCL}}\|_F \leq \exp\left(\frac{1}{2\tau^2 B} \sum_{t=0}^{T-1} \eta_t\right) \frac{1}{\tau\sqrt{B}} \left(\sum_{t=0}^{T-1} \eta_t\right) \Delta_{C,\delta}(B; \tau).$$

Combining the last two equations yields the lower bound on CKA_T with probability at least $1 - \delta$. \square

Corollary 2 (RSA lower bound). *In the setting of Thm. 1. Assume $\sigma_{D,T} > 0$. With probability at least $1 - \delta$,*

$$\text{RSA}_T \geq \frac{1 - r_T}{1 + r_T}, \quad r_T \leq \frac{\exp\left(\frac{1}{2\tau^2 B} \sum_{t=0}^{T-1} \eta_t\right) \frac{1}{\tau\sqrt{B}} \left(\sum_{t=0}^{T-1} \eta_t\right) \Delta_{C,\delta}(B; \tau)}{\sqrt{M} \sigma_{D,T}}.$$

Proof. Let $M = \binom{N}{2}$ and let $C := I - \frac{1}{M} \mathbf{1}\mathbf{1}^\top$ be the centering projector in \mathbb{R}^M . Write $a_c := Ca_T$ and $b_c := Cb_T$. Then

$$\text{RSA}_T = \frac{\langle a_c, b_c \rangle}{\|a_c\|_2 \|b_c\|_2}.$$

For any nonzero u and any v in an inner-product space,

$$\langle u, v \rangle = \langle u, u + (v - u) \rangle = \|u\|_2^2 + \langle u, v - u \rangle \geq \|u\|_2^2 - \|u\|_2 \|v - u\|_2,$$

and $\|v\|_2 \leq \|u\|_2 + \|v - u\|_2$. Therefore,

$$\frac{\langle u, v \rangle}{\|u\|_2 \|v\|_2} \geq \frac{1 - \|v - u\|_2/\|u\|_2}{1 + \|v - u\|_2/\|u\|_2}.$$

Apply this with $u = a_c$ and $v = b_c$ to obtain

$$\text{RSA}_T \geq \frac{1 - \|b_c - a_c\|_2/\|a_c\|_2}{1 + \|b_c - a_c\|_2/\|a_c\|_2}.$$

Since C is an orthogonal projector, $\|b_c - a_c\|_2 = \|C(b_T - a_T)\|_2 \leq \|b_T - a_T\|_2$. By construction of the RDM vectors,

$$b_T - a_T = -\text{vec}(\text{off}(\Sigma_T^{\text{NSCL}} - \Sigma_T^{\text{CL}})),$$

so $\|b_T - a_T\|_2 = \|\text{off}(\Sigma_T^{\text{NSCL}} - \Sigma_T^{\text{CL}})\|_F \leq \|\Sigma_T^{\text{NSCL}} - \Sigma_T^{\text{CL}}\|_F$. Finally, by Thm. 1, with probability at least $1 - \delta$,

$$\|\Sigma_T^{\text{NSCL}} - \Sigma_T^{\text{CL}}\|_F \leq \exp\left(\frac{1}{2\tau^2 B} \sum_{t=0}^{T-1} \eta_t\right) \frac{1}{\tau\sqrt{B}} \left(\sum_{t=0}^{T-1} \eta_t\right) \Delta_{C,\delta}(B; \tau).$$

Combining the last three displays yields the stated $(1 - r)/(1 + r)$ lower bound on RSA_T after substituting $\|a_c\|_2 = \sqrt{M} \sigma_{D,T}$. \square